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Q95Y35;
Q1-DEC-2001 (TrEMBLrel. 19, Created)
Q1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
Q1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Caenorhabditis elegans.
Eukaryota; Metazoa; Nemotoda; Chromadorea; Rhabditida; Rhabditoidea; Rhabditidae; Peloderinae; Caenorhabditis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL 38.2 KDA PROTEIN. Y108G3AL.2.

Q9CU16;
01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
1110021E09RIK PROTEIN (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (OCT-2001) to the ENEMBL; AC024744; AAK72068.1; --
Hypothetical protein.
SEQUENCE 327 AA; 38186 MW;
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MEDLINE-99069613;
Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S., Alzawa K., Izawa M., Nishi R., Kiyosawa H., Kondo S., Yamanaka I., Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R., Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T., Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H., Kuehl P., Lewis S., Matsuo Y., Nikaido I., Psole G., Quackenbush Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio Sakai K., Okido T., Furumo M., Aono H., Baldarelli R., Barsh G., Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (MAR-2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN-BRISTOL N2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            *Genome sequence of the nematode C. elegans: a investigating biology. The C. elegans Sequencin Science 282:2012-2018(1998).
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                                                                                                                         STRAIN-C57BL/6J; TISSUE-TESTIS;
MEDLINE-21085660; PubMed-11217851;
MEDLINE-21085660; PubMed-11217851;
MEDLINE-2108660; Shibata K.,
                                                                                                                                                        SEQUENCE FROM N.A
STRAIN-C57BL/6J; '
                                                                                                                                                                                                NCBI_TaxID=10090;
                                                                                                                                                                                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                        Mus musculus (Mouse).
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Rodentia;
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EMBL/GenBank/DDBJ databases
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Fukuda S.,
amanaka I.,
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Search completed: July 15, Job time: 1482 sec
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MGD; MGI:1914171; 1110021E09R1k.
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Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya
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01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                         closely related kinesin families."; J. Mol. Evol. 0:0-0(2001).
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Lawrence C.J., Malmberg R.L., Muszynski M.G., Da
"Maximum likelihood methods reveal conservation
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NCBI_TaxID=4577;
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Spermatophyta; Magnollophyta; Liliopsida; Poales; Poaceae; PACC cl
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Eukaryota; Viridiplantae;
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EMBL: AK015938; BAB30042.1;
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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Copyright (c) 1993 - 2000 Com
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AAB86295
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                                                                                                                                                                                                                                                                                     SUMMARIES
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VK005 VK region. 
Antibody variable 
Murine 1D9 antibod
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Mouse derived ligh
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AAR52058 AAR82961 AAR68741 AAR79248 AAR95217	AAM77590 AAR24808 AAR42803 AAR99849 AAY95225 AAY95227 AAE06948 AAE06948 AAE06996	AAM64423 AAM77235 AAM21171 AAM37382 AAM38594 AAW58494 AAY39809 AAR38159	AAE06961 AAE06962 AAE06963 AAE06963 AAE06965 AAE06965 AAE06966 AAE06967 AAE06967 AAE06967 AAE06967 AAE06967 AAE06967
Light chain variab Antibody 38193 lig MAD B3 light chain Light chain variab B3 immunoglobulin	Human bone marrow RSV19 VK. Mus mus RSV19 light chain Murine antibody 58 Anti-platelet glyc Anti-platelet glyc Anti-platelet glyc Human HF-21/28 kap Human kappa light Humanised Mab ligh		Mouse germline kap Fight chain variab Anti-DNA antibody Peptide #10995 enc

44409877114333344410987714441

ALIGNMENTS

RESULT AAW79207

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AAW79207 standard;

Protein;

32 \$

Framework 3 region of human V kappa gene HUM5400

21-DEC-1998 AAW79207;

(first entry)

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07-JUN-1995;
05-MAR-1993;
09-SEP-1993;
29-MAR-1995;
                                                                                                                                                                                           Monoclonal antibody; MAb; LO-CD2a; humanised antibody; CD2 antigen; human lymphocyte; immune response; chimeric; graft-versus-host disease; T-cell; transplant rejection; autoimmune disease; HUM5400.
                                                                                                                     07-JUN-1995;
                                                                                                                                       06-OCT-1998
                                                                                                                                                         US5817311-A
                                                                                                                                                                          Homo sapiens
                  WPI; 1998-556337/47.
                                     Bazin
                                                    (UYLO-) UNIV CATHOLIQUE LOUVAIN
Inhibition of T-cell mediated immune response with anti-CD2
                                    Ή,
                                    Latinne D;
                                                                        95US-0472281.
93US-0027008.
93US-0119032.
95US-0407009.
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germline

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This invention describes novel antibodies (Ab) in which the variable region (VR) of at least one chain and/or the VR of at least one heavy chain includes at least one of 7 specified sequences, or fragments of these sequences, or contain at least one light chain and/or heavy chain encoded by specific nucleic acid sequences (I) and (II), reproduced, or their fragments. The products of the invention have
                                                                                                 Claim
                                                                                                                                                                    WPI;
                                                                                                                             diagnosis
                                                                                                                                                                                            Mueller-Hermelink HK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human V kappa gene HUM5400. This is used to consruct a humanised antibod LO-CD2a. The invention relates to the use of the monoclonal antibody (MAB) LO-CD2a or a humanised or a chimeric version of the LO-CD2a antibody for the inhibition of a T-cell mediated lumune response in a patient. The MAB LO-CD2a (produced by hybridoma cell line AFCC HB 11423) can bind to an epitope on the CD2 antigen of the human lymphocytes. The T-cell mediated immune response in a patient can be inhibited by administering the MAB LO-CD2a and antibody that binds to the same administering the MAB LO-CD2a and antibody that binds to the same
                                                                                                                                                                                                                                                              23-DEC-1999;
                                                                                                                                                                                                                                                                                        23-DEC-1999;
                                                                                                                                                                                                                                                                                                                  28-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Murine derived antibody Wue-1 light chain variable region
                                                                                                                                                                                                                        (GREI/)
                                                                                                                                                                                                                                    (MUEL/)
                                                                                                                                                                                                                                                                                                                                                                                              tumor; multiple myeloma;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   diseases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         monoclonal antibody 10-CD2a - used for preventing transplant rejection or for treating graft-versus-host disease or auto-immune
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ftlkisr 21
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7; Conserv
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t
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                                                                                                                             autoimmune
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Pred. No.
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; plastocytoma; FW-3.
                                                                                                                                                                                                                                                                                                                                                                                                                       light chain; heavy chain;
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ma cell
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Matches 7
                   30-JUN-1994;
                                        27-JUN-1995;
                                                                06-MAR-1996
                                                                                      EP699755-A2
                                                                                                                                                                                                                                                                                                                                                                     variable region; epidermal growth factor receptor; hybridoma; framework; cloning; computer; algorithm; immunogenicity; site-directed mutagenesis; T-lymphocyte epitope; tertiary strpoint mutation; antibody engineering; protein engineering;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antitumor and immunomodulatory activity. Ab, or other antibodies that recognize the same antigen, are used: (1) to identify cognate antigens; (11) for specific labeling of plasma cells (PC), for identification or separation, e.g. in an extracorporeal system; (111) for generating additional antibodies able to label PC; and (1v) for treating autoimmune diseases and/or tumors, e.g. multiple myeloma, lymphoma and/or plastocytoma. Ab are specific for mature PC, i.e. they do not recognize precursor stages, even though these precursors are used as immunogens. As therapeutic agents, they should show fewer side effects than conventional chemotherapeutic agents. This sequence represents the Wue-1 antibody variable region light chain FW-3 fragment described in the method of the
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                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR92994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR92994
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Similarity 100
7; Conservative
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  12-OCT-1995
                                                    W09527045-A1.
                                                                                                                                                     Region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pattern
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The sequence represents residues from a human immunoglobulin with homology to the light chain variable region from monoclonal antibody IOR-R3 (AAR92993), specific for epidermal growth factor receptor, produced by a mouse hybridoma. The sequence is partial, and complementarity determining regions are omitted. The sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying interspecies differences in amino acid sequence of Ig
T-cell epitopes - by sequence comparison, also humanised antibodies
contg. altered T-cell epitopes, retaining antigen specificity but
                                                                                                                                                                                                    Region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Light chain; RT3; murine; catalytic antibody; bacteriophage;
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nes 7; Conserv
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43..49
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29..42
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                                                                                                                                 Treatment and preventive agent for hypercalcaemia - contg. one of anti-human para-thyroid-hormone-related protein monoclonal antibod anti-human chimera monoclonal antibody, fused gene and cell
                                                                Disclosure; Page 13; 18pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                   15-MAY-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JP04228089-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antihuman parathyroid hormone-related protein; monoclonal antibody; variable region; rodent/human chimeric MAb; constant region; PTHTP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hypercalcaemia
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                                                                                                                                                                                                                                    WPI; 1992-320987/39.
                                                                                                                                                                                                                                                                                   (KANF ) KANEKA CORP.
                                                                                                                                                                                                                                                                                                                                    15-MAY-1990;
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The sequences given in AAR27008-11 are fragments which were used the construction of an agent for the treating and prevention of

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RESULT

AAY56670

ID AAY5

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AC AAY5

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                                                                                                                                                                                                                                                              (complementarity determining regions) derived from a non-human antigen-
specific donor antibody, and an acceptor framework from a non-human
primate. The Abs are prepared by grafting CDRs from a non-human antigen-
specific donor antibody onto homologous Old World ape or monkey acceptor
frameworks. The Abs have reduced immunogenicity and are better tolerated
in humans (because of the close similarity between the human and primate
principles), but retain the full antigen-binding affinity of the donor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Complementarity determining region; antibody; primate; immunogenicity; Old World ape; Old World monkey; antigen-binding affinity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antibodies containing donor complementarity determining regions and non-human primate acceptor frameworks, having reduced immunogenicity humans -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 26; Page 101; 123pp; English
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                      AAR72066;
                                                                   AAR72066 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                              gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 66; 94pp; English.
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Best Local S
Matches 7
                                                                    Tumour-associated antigen mucin-1; MUC-1; adenocarcinoma; VL; variable light chain region; cancer; breast; ovary; lung; bladder, cytostatic; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tissue and clones encoding autoimmune-associated immunoglobulin fragments were obtained. 14/15 clones of L chain (Kappa) regions showed homology to the putative vK germline gene Kull2 (given in AAQ89317) and the remaining clone, OFTK.7 (AAQ89326), to the VKOO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Graves' ophthalmopathy-associated monoclonal antibody - produced by molecular cloning of immunoglobulin genes by \ensuremath{\mathsf{PCR}}
                           Unidentified
                                                                                                                                                                      Antibody variable light chain region (VL)
                                                                                                                                                                                                                         04-JAN-2002
                                                                                                                                                                                                                                                                             AAE12711;
                                                                                                                                                                                                                                                                                                                         AAE12711 standard; peptide; 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                         (first entry)
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                                                                                                                                           The invention relates to an isolated tumour-associated antigen mucin-1 (MUC-1)-specific binding member comprising an antigen binding domain cregion having an antibody variable light (VL) or heavy (VH) region, and the properties of the properties of the properties of the binding member is useful for diagnosing cancer, preferably adenocarcinoma the binding of MUC1-specific binding member to MUC1 is detected by a detection method selected from enzyme-linked immunosorbent assay, and the binding terminal properties of the containing cancer, preferably competed to the containing cancer preferably contained imaging, scintillation counting, and x-ray film. MUC1-specific binding member is useful for treating cancer, preferably containing cancer, in an individual, where the cancer is present in tissue of the breast, ovary, lung, or bladder of the individual. MUC1-specific binding member is useful for diagnosing and imaging MUC1-expressing cancer cells and tissues, for purifying or isolating non-glycosylated, containing molecules, and for therapeutically or prophylactically containing molecules, and for therapeutically or prophylactically.
                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel isolated tumor-associated antigen mucin-1-specific binding member for diagnosing and treating cancer, comprises mucin-1 binding domain or its portion for binding to an epitope of the protein core of mucin-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hoogenboom HRJM,
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1 FTLKISR 7
                                                                                                                                                                           19 cancer. The present sequence is antibody variable light chain
(VL) from DP47 germ line.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 35; 126pp; English.
                                                                                                                                         100 AA;
                                       Conservative
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                                                        Score 33;
Pred. No.
                                       Mismatches
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RESULT 1
AAE06958
ID AAEC 밁 Murine 1D9 AAE06958; AAE06958 standard; Protein; 100 16-OCT-2001 76 ftlkisr 82 10 antibody partial kappa light chain variable (VK) region. (first entry)

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anaphylaxis; malignancy, inflammation; stenosis; allograft rejection; fibrotic disease; angioplasty; acquired immune defictency syndrome; AIDS; inflammatory glomerulopathy; vascular intervention; 1D9 antibody; neointimal hyperplasia; VK; kappa light chain variable region. Mus sp multiple sclerosis; atherogenesis; atherosclerosis; restenosis; asthma; HIV infection; cytostatic; vasotropic; leukocyte trafficking; allergy; inflammatory disorder; autoimmune disorder; rheumatoid arthritis; shoc Murine; humanised antibody; CC-chemokine receptor 2; CCR2; nephrotropic; neuroprotective; immunosuppressive; human immunodeficiency virus;

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                                                                                                                                                                                                                                                                                                  RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The patent discloses a humanised antibody or its antigen-binding CC Crimmokine receptor 2 CC (CR20), comprising an antigen binding region of non-human origin. The CC and at least a portion of an immunoglobulin of human origin. The CC and at least a portion of an immunoglobulin of human origin. The CC humanised antibodies are useful for inhibiting or treating CC HIV infection. The proteins of the invention are useful for inhibiting or treating CC leukocyte trafficking, for treating CCR2-mediated disorders such as cCC inflammatory disorder, autoimmune disorders such as rheumatold arthritis and multiple sclerosis, atherogenesis and atherosclerosis, CC and in the manufacture of a medicament for treating CCR2-mediated disgnosis, CC and for inhibiting restenosis. They are useful in therapy or diagnosis, CC and for inhibiting restenosis. They are useful in therapy or diagnosis, CC and for inhibiting restenosis, They are useful in therapy or diagnosis, CC and for inhibiting restenosis, They are useful for treating CCR2-mediated dispancy, chronic and acute inflammation, histmanic and Tg2-CC mediated allergic reaction, shock, stenosis, allograft rejection, CC fibrotic disease, sathma, inflammatory glomerulopathies, acquired CC intervention, including angioplasty and/or stent placement in a mammal. CC intervention, including angioplasty and/or stent placement in a mammal. The second of a vessel in a mammal, proferably associated with vascular intervention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
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                     inflammatory disorder; autoimmune disorder; rheumatoid arthritis; shock; multiple sclerosis; atherogenesis; atherosclerosis; restenosis; asthma; anaphylaxis; malignancy; inflammation; stenosis; allograft rejection;
                                                                           HIV infection; cytostatic; vasotropic; leukocyte trafficking; allergy;
                                                                                            Mouse; humanised antibody; CC-chemokine receptor 2; CCR2; nephrot neuroprotective; immunosuppressive; human immunodeficiency virus;
                                                                                                                                                      Mouse
                                                                                                                                                                                         16-OCT-2001
                                                                                                                                                                                                                             AAE06959
                                                                                                                                                                                                                                                                AAE06959 standard; Protein;
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tibrotic
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                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
les 7; Conser
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                                                                                                                                                                                                                                                                                                                                                       ftlkisr 82
                                                                                                                                                  germline kappa light chain variable (VK) region,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence is murine 109 antibody partial kappa light chain
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                                                                                                                                                                                       (first entry)
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     angioplasty;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 33;
Pred. No.
     acquired immune
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       deficiency syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 100;
                                                                                                              CCR2; nephrotropic
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(CCR2), comprising an antigen binding region of human origin. The and at least a portion of an immunoglobulin of human origin. The humanised antibodies are useful for inhibiting the interaction of a cell expressing CCR2. They are useful for inhibiting or treating HTV infection. The proteins of the invention are useful for inhibiting leukocyte trafficking, for treating CCR2-mediated disorders such as the unatold
                                                                              malignancy, chronic and acute inflammation, histamine and IgE-
mediated allergic reaction, shock, stenosis, allograft rejection,
fibrotic disease, asthma, inflammatory glomerulopathies, acquired
immune deficiency syndrome (AIDS), restenosis associated with vascular
intervention, including angioplasty and/or stent placement in a mammal.
Humanised antibodies are also useful for inhibiting narrowing of the
lumen of a vessel in a mammal, and inhibiting neointimal hyperplasia of
a vessel in a mammal, preferably associated with vascular intervention.
                                                                                                                                                                                                                                 inflammatory disorder, autoimmune disorders such as rheumatoid arthritis and multiple sclerosis, atherogenesis and atherosclerosis, and for inhibiting restenosis. They are useful in therapy or diagnosis, and in the manufacture of a medicament for treating CCR-2 mediated disease. They are also useful for treating allergy, anaphylaxis, mallgnancy, chronic and acute inflammation, histamine and IgE-mallgnancy, chronic and acute inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            The patent discloses a humanised antibody or its antigen-binding fragment, having binding specificity for CC-chemokine receptor 2 (CCR2), comprising an antigen binding region of non-human origin and at least a portion of an immunoglobulin of human origin. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Humanized immunoglobulin for treating a CC-chemokine receptor 2-mediated disorder in a patient, comprises a binding specificity
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                                                                The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 147; 183pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCR2,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and a non-human antigen binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Horvath C,
                                          70/3.
                                                                mouse germline
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                                                            RESULT
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                                                                                                               Best Local Similarity
                                     AAE06960
                                                AAE06960 standard; Protein; 100
                                                                                76
                                                             12
                                                                                            1 FTLKISR 7
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                                                                                                         Conservative
                                                                                                              100.0%;
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                                                                                                              Score 33;
Pred. No.
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                                                                                                         Mismatches
                                                                                                                     DB
                                                                                                                    22;
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                                                                                                                     Length 100;
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Sequence

100 AA;

Mouse;

16-OCT-2001

(first

entry)

Mouse germline kappa light chain variable (VK) region, 70/1.

humanised antibody; CC-chemokine receptor 2; CCR2; nephrotropic;

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RESULT
AAE06961
ID AAE
XX
AC AAE
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                                                                                                                                                                                                                                                                                                                                                                                                                                            The patent discloses a humanised antibody or its antigen-binding CC cfragment, having binding specificity for CC-chemokine receptor 2 CC (CCR2), comprising an antigen binding region of non-human origin. The cand at least a portion of an immunoglobulin of human origin. The chamanised antibodies are useful for inhibiting the interaction of a cell expressing CCR2. They are useful for inhibiting or treating CC leukocyte trafficking, for treating CCR2-mediated disorders such as cleavers and multiple sclerosis, atherogenesis and atherosclerosis, and in the manufacture of a mediatement for treating CCR2-mediated disorders such as rheumatoid cand in the manufacture of a medicament for treating CCR2-mediated disorders. They are also useful in therapy or disposis, and in the manufacture of a medicament for treating CCR2-mediated disease. They are also useful for treating cCR2-mediated disease. They are also useful for treating cory anaphylaxis, canding an antigent of the mediated allergic reaction, shock, stenosis, allograft rejection, cory and acute inflammatory, histamine and Lgc mediated allergic reaction, shock, stenosis, allograft rejection, cory and the second a
                                                                                                                                                                                                                                                                                    Query Match
Best Local
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                                                AAE06961 standard; Protein; 100 AA
                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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HTV infection; cytostatic; vasotropic; leukocyte trafficking; allergy;
inflammatory disorder; autoimmune disorder; rheumatoid arthritis; shock;
multiple scierosis; atherogenesis; atherosclerosis; restenosis; asthma;
                                                                                                                                                                      76
                                                                                                                                                                                                                                                             Local Similarity nes 7; Conserv
                                                                                              13
                                                                                                                                                                                                               1 FTLKISR 7
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                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                    Score 33;
Pred. No.
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                                                                                                                                                                                                                                                                                                           Length 100;
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inflammatory disorder, autoimmune disorders such as rheumatoid arthritis and multiple sclerosis, atherogenesis and atherosclerosis, and for inhibiting restenosis. They are useful in therapy or diagnosis, and in the manufacture of a medicament for treating CCR-2 mediated disease. They are also useful for treating allergy, anaphylaxis, malignancy, chronic and acute inflammation, histmaine and IgE-mediated allergic reaction, shock, stenosis, allograft rejection, fibroric disease, astima, inflammatory glomerulopathies, acquired immune deficiency syndrome (AIDS), restenosis associated with vascular intervention, including angioplasty and/or stent placement in a mammal. Humanised antibodies are also useful for inhibiting nearrowing of the lumen of a vessel in a mammal, and inhibiting neointimal hyperplasia of a vessel in a mammal, preferably associated with vascular intervention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     fibrotic disease; angioplasty; acquired immune deficiency syndrome; AIDS; inflammatory glomerulopathy; vascular intervention; neointimal hyperplasia; VK; kappa light chain variable region.
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anaphylaxis; malignancy; inflammation; stenosis; allograft rejection;
                                                                                                                                                                                                                                                                                                                                                                     a cell expressing CCR2. They are useful for inhibiting or treating HIV infection. The proteins of the invention are useful for inhibiting leukocyte trafficking, for treating CCR2-mediated disorders such as inflammatory disorders such inflammatory disorders and increases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The patent discloses a humanised antibody or its antigen-binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Humanized immunoglobulin for treating a CC-chemokine receptor 2-mediated disorder in a patient, comprises a binding specificity for CCR2, and a non-human antigen binding region and human immunoglobulin
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                                      The present sequence is (VK) region, 70/2.
                                                                                                                                                                                                                                                                                                                                                                                                                                              fragment, having binding specificity for CC-chemokine receptor 2 (CCR2), comprising an antigen binding region of non-human origin and at least a portion of an immunoglobulin of human origin. The humanised antibodies are useful for inhibiting the interaction of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 148; 183pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /label= Unknown
                                                            mouse germline kappa light chain
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Query Match
Best Local Similarity
"~+~hes 7; Conserv

Conservative

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100.0%;

Score 33; Pred. No. Mismatches

DB 22;

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                                                                                                                  lumen of a vessel in a mammal, and inhibiting neointimal hyperplasia of a vessel in a mammal, preferably associated with vascular intervention. The present sequence is mouse germline kappa light chain variable
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            patent discloses a humanised antibody or its antigen-binding
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Best Local Similarity
Thes 7; Conserve
                                                                                                                                                                                                                                                               inflammatory disorder, autoimmune disorders such as rheumatoid arthritis and multiple sclerosis, atherogenesis and atherosclerosis, and for inhibiting restenosis. They are useful in therapy or diagnosis, and in the manufacture of a medicament for treating CCR-2 mediated disease. They are also useful for treating allergy, anaphylaxis, malignator, abroats is figure to be the property of the state of the sta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The patent discloses a humanised antibody or its antigen-binding fragment, having binding specificity for CC-chemokine receptor 2 (CCR2), comprising an antigen binding region of non-human origin and at least a portion of an immunoglobulin of human origin. The humanised antibodies are useful for inhibiting the interaction of a cell expressing CCR2. They are useful for inhibiting or treating HIV infection. The proteins of the invention are useful for inhibiting leukocyte trafficking, for treating CCR2-mediated disorders such as information and the contraction are useful for inhibiting contractions.
                                                                                   mediated allergic reaction, shock, stenosis, allograft rejection, fibrotic disease, asthma, inflammatory glomerulopathles, acquired immune deficiency syndrome (AIDS), restenosis associated with vascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Humanized immunoglobulin for treating a CC-chemokine receptor 2-mediated disorder in a patient, comprises a binding specificity for CCR2, and a non-human antigen binding region and human immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Humanized immunoglobulin 2-mediated disorder in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  anaphylaxis; malignancy; inflammation; stenosis; allograft rejection; fibrotic disease; angioplasty; acquired immune deficiency syndrome; AIDS; inflammatory glomerulopathy; vascular intervention;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neuroprotective; immunosuppressive; human immunodeficiency virus; HIV infection; cytostatic; vasotropic; leukocyte trafficking; allergy; inflammatory disorder; autoimmune disorder; rheumatoid arthritis; shock
                                                                                                                                                                                                                   malignancy, chronic and acute inflammation, histamine and IgE-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 149; 183pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-4888888/53
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    antibodies are also
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                                            including angioplasty and/or stent placement in
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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CC lumen of a vessel in a mammal, and inhibiting neointimal hyperplasia of CC a vessel in a mammal, preferably associated with vascular intervention. CC The present sequence is mouse germline kappa light chain variable CC (VK) region, V-IC.

XX Sequence 100 AA:

Query Match
Best Local Similarity 100.0%; Score 33; DB 22; Length 100;
Best Local Similarity 100.0%; Pred. No. 4;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 FTLKISR 7

Db 76 ftlkisr 82

Search completed: July 15, 2002, 12:57:58

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are greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            July 15, 2002, 12:59:18; Search time 75.67 Seconds (without alignments) 2.260 Million cell updates/sec
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Copyright (c) 1993 - 2000 Comp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/BCTUS_COMB.pep:*
/cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
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US-07-297-696C-71

US-08-12-930B-73

US-08-472-281A-33

US-08-472-281A-33

US-08-976-288A-71

US-08-497-312-19

US-08-481-133-151

US-08-481-133-151

US-08-477-484B-151

US-08-477-788A-151

US-08-477-788A-151

US-08-477-788A-151

US-08-477-531B-15

US-08-477-531B-15

US-08-478-201-151

US-08-378-939-28

US-08-378-939-28

US-08-378-939-28

US-08-389-765-151

US-08-389-765-151

US-08-389-765-151

US-08-389-765-151

US-08-381-398-48

US-08-053-171-15

US-08-053-171-15

US-08-053-171-15

US-08-053-171-16

US-08-053-171-16

US-08-053-171-16

US-08-053-171-16
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  Sequence
33, Appl

71, Appl

33, Appl

33, Appl

33, Appl

34, Appl

37, Appl

37, Appl

37, Appl

38, Appl

15, Appl

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Result No.

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US-08-484-508-18	US-08-752-844-15	US-08-678-194-6	US-08-606-293-8	US-08-606-293-4	US-08-859-649-29	US-08-859-649-19	US-08-483-199-18	US-08-472-281A-89	US-08-475-000-18	US-08-388-672A-25	US-08-476-349A-89	US-08-476-349A-88	US-08-077-252B-3	US-08-477-877B-89	US-08-478-039-89	US-08-478-039-88	US-07-942-245-28
Sequence 18, Appl	Sequence 15, Appl	Sequence 6, Appli	Sequence 8, Appli	Sequence 4, Appli	Sequence 29, Appl	•	Sequence 18, Appl	`	•	Sequence 25, Appl	Sequence 89, Appl	Sequence 88, Appl	Sequence 3, Appli	Sequence 89, Appl	Sequence 89, Appl	Sequence 88, Appl	Sequence 28, Appl

ALIGNMENTS

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; TOPOLOGY: 1:
; MOLECULE TYPE:
US-08-477-877B-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Bazin, Herv
APPLICANT: Latinne, Dominique
TITLE OF INVENTION: LO-CD2a Antibody and Uses Thereof for Inhibiting T-Cell Act
NUMBER OF SEQUENCES: 96
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
                                                                                                                                             TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/477, E
FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/407,009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 6 DEC.
STREET: ROSeland
STATE: New Jersey
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/0:
FILING DATE: 05-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Olstein, Elliot M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inc
COMPUTER: IBM PS/2
                                                                                                                                                                                      REGISTRATION NUMBER: 24,025
REFERENCE/DOCKET NUMBER: 61
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
                                                                                                                           SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/4
FILING DATE: 29-MAR-1995
APPLICATION NUMBER: 08/1
FILING DATE: 09-SEP-1993
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                                                            STRANDEDNESS:
                                                                                                     LENGTH:
                                                                                 amino acid
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EE: Cecchi, Stewart & Olstein 
6 Becker Farm Road
                                                                                                       32 amino acids
                                             linear
                    polypeptide
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Query Match
Best Local Similarity
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                                                                             US-08-129-930B-71

: Sequence 71, Application US/08129930B

: Patent No. 5804187
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US-07-977-696C-71
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Best Local
                                                                 GENERAL INFORMATION:
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             APPLICANT:
APPLICANT:
APPLICANT:
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   APPLICANT:
                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS
LENGTH: 32 amino acid:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 30,930
REFERENCE/DOCKET NUMBER: P6
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 748-6868
TELEFAX: (510) 748-6688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS 5.0

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA: 805/07/977,696C

FILING DATE: 11-16-92

CLASSIFICATION: 435

CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Analogue Peptides with SITILE OF INVENTION: for Carcinomas and Kit at TITLE OF INVENTION: and Therapeutic Methods.
NUMBER OF SEQUENCES: 81
CORRESSIONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Amzel Ph.D., Viviana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: do Couto, remando ....
APPLICANT: Ceriani Dr., Roberto L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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| 15 FTLKISR 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: PRETTY, SCHROEDER & POPLAWSKI STREET: 444 South Flower Street, Suite 2000 CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
nes 7; Conserv
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o. 5792852
                                                                                                                                                                                                                                                                                                                                                                     amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             California
do Couto Dr., Fernando .
Cerlani Dr., Roberto L.
Peterson Dr., Jerry A.
Padlan Dr., Eduardo A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
                                                                                                                                                                                                                                              Conservative
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                                              Fernando J.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Jerry A.
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Pred. No. 0.72;
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Pred. No. 0.72;
                                                                                                                                                                                                                                           Mismatches
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US-08-472-281A-33
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                                                                                                                                                                                                                                                                                                                                                                                              Patent No. 581731
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 33, Application US/08472281A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                               OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inc
                                                                                                                                                                                                                                                                                                        APPLICANT: Bazin, Herv
APPLICANT: Latione, Dominique
APPLICANT: LO-CD2a Antibody and Uses Thereof for Inhibiting T-Cell Act
TITLE OF INVENTION: LO-CD2a Antibody and Uses Thereof for Inhibiting T-Cell Act
                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: AMZEI Ph.D., VIVIANA REGISTATION NUMBER: 30,930 REFERENCE/DOCKET NUMBER: CR. TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 2022
CITY: Walnut Creek
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
APPLICATION NUMBER: US/08/472,281A FILING DATE: 07-JUN-1995 CLASSIFICATION: 424
                                                                                                          COMPUTER:
                                                                                                                                                            ZIP: 07068
                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                STATE:
                                                                                                                                                                                                            CITY: Roseland
                                                                                                                                                                                                                                                ADDRESSEE: Carella, Byrne, Bain, Gilfillan, ADDRESSEE: Cecchi, Stewart & Olstein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15 FTLKISR 21
                                                                                                                                                                                                                                 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity tes 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/129,930B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
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                                                                                                                                                                          New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               32 amino acids
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2055 No. 5804187th Broadway, Suite 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2: (510) 521-1333
(510) 521-3541
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                       IBM PS/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IBM PC compatible
SYSTEM: PC-DOS/MS-DOS 5.0
PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
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                                                                                                                          inch diskette
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Carcinoma Specificity, and K:
Diagnostic Vaccination and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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Pred. No. 0.72;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 32;
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Query Match
Best Local Similarity
Watches 7; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Postema, Christina E.
APPLICANT: White-Scharf, Mary
TITLE OF INVENTION: LO-CDZa Antibody and Uses
TITLE OF INVENTION: Thereof for Inhibiting
TITLE OF INVENTION: T-Cell Activation and
TITLE OF INVENTION: Proliferation
                                                                                          APPLICATION NUMBER: US/08/477,989B
FILING DATE: 07-UUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette
COMPUTER: IBM PS/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Olstein, Elliot M.
REGISTRATION NUMBER: 24,025
REFERENCE/DOCKET NUMBER: 61750-142
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 08/027,008
FILING DATE: 05 MAR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                              CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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APPLICATION NUMBER:
'FILING DATE: 29-MAR-
                       APPLICATION NUMBER: 08/11
APPLICATION NUMBER: 08/11
FTI.TNG DATE: 09-SEP-1993
                                                            APPLICATION NUMBER: 08/4
FILING DATE: 29-MAR-1995
                                                                                                                                                                              OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                   CITY: Roseland
STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                   STREET:
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                                                                                                                                                                                                                                                                    ZIP: 07068
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              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: (FILING DATE: 09-SEP-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                6 Becker Farm Road
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Cecchi, Stewart & Olstein
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05-MAR-1993
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                                             08/119,032
                                                                              08/407,009
            08/027,008
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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US-08-976-288A-71
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Best Local Similarity
""" Conserve
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APPLICANT: do Couto
APPLICANT: Ceriant
APPLICANT: Peterson
APPLICANT: Padlan I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 71, Application US/08976288A
                                 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: FITTLE OF INVENTION: CITILE OF INVENTION: IT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
TOPOLOGY: lir
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                 REFERENCE/DOCKET NUMBER: P6639938
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 622-7700
                                                                                                                                   APPLICATION NUMBER: 07/977,696
FILING DATE: NO. 6315997ember 16, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Viviana Amzel Ph.D.
REGISTRATION NUMBER: 30,930
                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                      FILING DATE: September 30, 1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                         CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 32 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: 201-994-1700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Olscein, Ellict M.
REGISTRATION NUMBER: 24,025
REFERENCE/DOCKET NUMBER: 61'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                    TELEFAX:
                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/976,288A
FILING DATE: No. 6315997ember 21, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 201-994-1744
                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: PC-DOS/MS-DOS 5.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 32 ....
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California
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32 amino acids
                                                                    (213) 489-4210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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Ceriani Dr., Roberto L.
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                                                                                                                                                                                                                                                                                                                                                                                            compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Analogue Peptides With Broad Carcinoma Specificity, and K Diagnostic Vaccination and
                                                                                                                                                                                                                                                                                                                                                            Release #1.0, Version #1.25
                                                                                                                                                                                                                                                         08/129,930
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Pred. No. 0.72;
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RESULT 8
US-08-881-037-37
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                                                                                                                        Query Match
Best Local Similarity
Tatches 7; Conserve
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: U5/08/497,312
FILING DATE: 30-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: CU 80/94
FILING DATE: 30-JUN-1994
ATTORNET/AGENT INFORMATION:
NAME: BOND, LAURENCE B.
REGISTRATION NUMBER: 2629US
REFERENCE/DOCKET NUMBER: 2629US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 801/532-1922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 801/531-9168
TELEX: 388961 1PM04UT
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                        TOPOLOGY: un
MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Method for obtaining mod
TITLE OF INVENTION: immunoglobulins with red
TITLE OF INVENTION: antibody variable domain:
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: CENTRO DE INMUNOLOGIA MOLECULAR
STREET: 215 Y 15, ATABEY PLAYA
                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 81 amino acids
TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 601/531-9168
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STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15 FTLKISR 21
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                                                                     53 FTLKISK 59
                                                                                                                                                                                                                                                                                           STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 11600
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                                                                                                      1 FTLKISR 7
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5712120
                                                                                                                                     100.0%; Score 33; 1 ilarity 100.0%; Pred. No. 1. Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                             unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Method for obtaining modified immunoglobulins with reduced immunogenicity of murine antibody variable domains, compositions containing them.
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                                                                                                                                                                        Length 81
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US-08-488-113B-151

: Sequence 151, Application US/08488113B

: Patent No. 5744580
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                                                                                                                                                                                                                                                                                                                                                                              RESULT
                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Better, Marc D.
APPLICANT: Carroll, Stephen F.
APPLICANT: Studnika, Gary M.
TITLE OF INVENTION: Immunotoxins
TITLE OF INVENTION: Proteins
NUMBER OF SEQUENCES: 169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 37, Application US/08881037 Patent No. 6080588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/881,037
APPLICATION NUMBER: US/08/881,037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 104 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER: 34,202
REFERENCE/FOCKET NUMBER: 07
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650) 813-5500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Konski, Antoinette F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
FILING DATE: 18-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison & Foerster
STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Glick, Gary D.
APPLICANT: Swanson, Patrick C.
TITLE OF INVENTION: DNA BINDING ANTIBODIES
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 94304-1018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 755 Page
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: sin
                                        ZIP: 60661
                                                                            CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                     68 FTLKISR 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (650) 494-0792
                                                                                                                    STREET:
                                                                                                                  ADDRESSEE: McAndrews, Held & Malloy, Ltd.
STREET: 500 West Madison Street, 34th floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 FTLKISR 7
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                                                             USA
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ilarity 100.0%;
Conservative 0
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                                                                                                                                                                                                                   Immunotoxins Comprising Ribosome-Inactivating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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CURRENT APPLICATION DATA:

PatentIn Release #1.0, Version #1.25

APPLICATION NUMBER:

US/08/488,113B 07-JUN-1995 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS

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Query Match
Best Local Similarity
"""+" 7; Conserve
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US-08-477-484B-151
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US-08-488-113B-151
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                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 108 amino acids
TYPE: amino acid
             SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                               TITLE OF INVENTION: Immunoto:
                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                            APPLICANT: Studnika, Gary M.
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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 0
FILING DATE: 04-NOV-1991
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 1
FILING DATE: 09-DEC-
PRIOR APPLICATION DATA:
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APPLICATION NUMBER:
FILING DATE: 12-MAY-
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                                                                                                                        COUNTRY:
                                                                                                                                          STATE:
                                                                                                                                                      CITY: Chicago
                                                                                                                                                                     STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                     72 FTLKISR 78
APPLICATION NUMBER:
                                                                                                                                                                                 ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 07/901,707 FILING DATE: 19-JUN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 FTLKISR 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: McNicholas, Janet M.
REGISTRATION NUMBER: 32,918
                                                                                                          60661
                                                                                                                                       Illinois
                                                                                                                                                                E: McAndrews, Held & Malloy, Ltd.
500 West Madison Street, 34th floor
                                                                                                                        USA
                                                                                                                                                                                                                                                                           Better, Marc D.
Carroll, Stephen F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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09-DEC-1992
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                                                                                                                                                                                                                                           Immunotoxins Comprising Ribosome-Inactivating
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 US/08/477,484B
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Pred. No. 2
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Best Local Similarity
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              APPLICATION NUMBER: PCT/
FILING DATE: 14-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 32,918
REFERENCE/DOCKET NUMBER: 11022US07/200-70.P3.C2A
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/707-955
TELEX: 312/707-955
TELEX: 650 788-772
                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                            CLASSIFICATION: 424
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                               COMPUTER READABLE FORM:
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ATTORNEY/AGENT INFORMATION:
NAME: MCNicholas, Janet M
                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
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APPLICATION NUMBER: U
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                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
TOPOLOGY: 11
                                                                                                                                                                                                                                                   COUNTRY: United States of America ZIP: 20005-3934
                                                                                                                                                                                                                                                                                CITY: Washington
STATE: D.C.
                                                                                                                                                                                                                                                                                                                   STREET:
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                                                                                                                               APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 FTLKISR 7
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                                                                                                                                                                                                                                                                                                             E: Sterne, Kessler,
1100 New York Ave.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        108 amino acids
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                                                                                                             13-AUG-1993
13-DEC-1991
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09-DEC-1992
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12-MAY-1993
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                                                                                                                                                                                                                                                                                                                                                                  ka, Gary M.
Modified Antibody Variable Domains (as amended)
: 67
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                                                             PCT/US92/10906
                                                                                                                             us/08/107,669D
            US 07/808,464
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                                                                                                                                                                                                                                                                                                             Goldstein and Fox P.L.L.C
N.W., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 108;
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ATTORNEY/AGENT INFORMATION:

Michele

Cimba

33,851

0610.1000001/MAC

REGISTRATION NUMBER: 33,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-472-788A-15
; MOLECULE TYPE: protein US-08-472-788A-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15, Application US/08472788A Patent No. 5770196 GENERAL INFORMATION:
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ELENGTH: 108 amin Type
                                                                                                            INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/808,464
FILING DATE: 13-DEC-1991
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/472,788A
FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Modified Antibody Variable Domains NUMBER OF SEQUENCES: 89
CORRESPONDENCE ADDRESS:
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TELEPHONE: 202/371-2600
                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 108 amino acid
                                                                                                                                             REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 0610.1000003
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202/371-2600
TELEFAX: 202/371-2540
                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 08/082,842 FILING DATE: 23-JUN-1993 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 424
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: LLUU New
CITY: Washington
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nes 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Sterne, Kessler, Goldstein and Fox P.L.L.C
STREET: 1100 New York Ave., N.W., Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             72 FTLKISR 78
                                    TOPOLOGY: linear
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                                                Query Match
Best Local Similarity
Thes 7; Conservi
                                                                                                                                            ; MOLECULE TYPE: protein US-08-477-5318-15
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                              TELEFAX: 202/371-2540
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPUTED:
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 07/808,464
FILING DATE: 13-DEC-1991
ATTORNEY/AGENT INFORMATION:
NAME: Michele A. Cimbala
                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 08/107,669
APPLICATION NUMBER: 08/107,669
FILING DATE: 13-AUG-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/10906
FILING DATE: 14-DEC-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: 202/371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Studnicka, TITLE OF INVENTION: MC
                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 07 CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 0610.1000004/MAC
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72 FTLKISR 78
                                                                                                                                                                              TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                             TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/477,531B FILING DATE: 07-JUN-1995
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                                1 FTLKISR 7
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o. 5821123
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1100 New York Ave., N.W., Suite 600
                                                                                                                                                                                                              108 amino acids
                                                                                                                                                                                                                                                                202/371-2540
                                                                 Conservative
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Modified Antibody Variable Domains (as amended)
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Pred. No. 2.3;
                                                                              Score 33; DB
Pred No. 2.3;
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                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Version #1.25
                                                                                                DB 2;
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                                                                 Gaps
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RESULT 14 US-08-646-360-151 ; Sequence 151, Application US/08646360 ; Patent No. 5837491

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WESULT 15
US-08-082-842A-15
Sequence 15, Application US/08082842A
Patent No. 5869619
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Best Local Similarity
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                GENERAL INFORMATION:
APPLICANT: Studnicka, Gary M.
TITLE OF INVENTION: Modified Antibody Variable Domains
NUMBER OF SEQUENCES: 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: 650 388-1248
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARB: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REPERENCE/DOCKET NUMBER: 200-70.P4
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/707-8889
TELEPAX: 312/707-9155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/787,567
FILING DATE: 04-NOV-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 08/064,691
FILING DATE: 12-MAY-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/988,430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: PCT/
FILING DATE: 12-MAY-1994
PRIOR APPLICATION DATA:
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PRIOR APPLICATION NUMBER: PCT/US94/05348
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TITLE OF INVENTION: Proteins
NUMBER OF SEQUENCES: 173
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/901,707
FILING DATE: 19-JUN-1992
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TOPOLOGY: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: McNicholas, Janet M. REGISTRATION NUMBER: 32,918
                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                                                                                                                                                                                                                       1 FTLKISR 7
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500 West Madison Street, 34th floor
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Carroll, Stephen F.
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Search completed: July 15, 2002, 12:59:19 Job time: 390 sec
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Best Local Similarity
Wetches 7; Conserve
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                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 33,851
REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202/371-2600
TELEPAX: 202/371-2540
TRILEPAX: 202/371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 07
FILING DATE: 13-DEC-1991
ATTORNEY/AGENT INFORMATION:
NAME: Cimbala, Michele A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 14-DEC-1992 PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                      72
                                                                                                                                                                                                                                                                                                    TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                       1 FTLKISR 7
                                                                                                                                                                                                                                                                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: PCT/US92/10906
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                                                                                      FTLKISR 78
                                                                                                                                                                                                                                                                                              amino acid
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Conservative (
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Minimum
Maximum
                                                                                                                                                                                                                                                                                       Total number of hits satisfying chosen parameters:
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Perfect score:
                                                                                                                                                                       Result
                                                                                                                                                                                                                                                                                                                   Scoring table:
                                                                                                                                                                                                                                                                                                                                                                         OM protein -
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 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 2000000000
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Match Length
 PIR_71:*
1: pir1:*
2: pir2:*
3: pir3:*
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Gapop 10.0 , Gapext
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Listing first
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(c) 1993 - 2000 Comp
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S19974	PH1083	PH1067	рн1066	A42575	S45441	G38601	PH1070	PH0867	PH0863	S20648	S37504	S37506	S37522	S37516	S37532
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ALIGNMENTS

Ig kappa chain V-III region (CP1) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 23-Oct-1992 #sequence_revision 23-Oct-1992 #text_change 09-May-1997
C;Accession: JT0521

patients with x-linked agammagl

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chain

rearrangeme

RESULT JT0521

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R;Anker, R; Conley, M.E.; Pollok, B.A.
J. Exp. Med. 169, 2109-2119, 1989
A;Title: Clonal diversity in the B cell repertoire of patients wi A; Reference number: JT0511; MUID:89279157
A; Accession: JT0521
A; Molecule type: mRNA
A; Residues: 1-54 <ANK>
A; Note: the sequence shown here is one of eight productive V-D-J: C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Reywords: heterotetramer; immunoglobulin
C; F:1-49/Domain: V region <VRE>
F; 10-54/Domain: J region <VRE>
                                                                                                                                                                                                                                                                                                                                                               Ig kappa chain V region - human (fragment)
(;Species: Homo sapiens (man)
(;Date: 20-Feb-1995 #sequence_revision 25-Oct-1996 #text_change 23-Jul-1999
(;Accession: S34082; S21526
R;Wagner, S.D.; Luzzatto, L.
Rr;Wagner, S.D.; Luzzatto, L.
Rur. J. Immunol. 23, 391-397, 1993
A;Title: V-Kappa gene segments rearranged in chronic lymphocytic leukemia are distrib
A;Reference number: S34076; MUID:93170387
A;Accession: S34082
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-71 <-WA2>

                                                                                                                                                                                                                              A;Cross-references: EMBL:X66042; NID:g33318; PIDN:CAA46841.1; PID:g33319 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin
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Best Local Similarity
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    Query Match
Best Local s
Matches 9
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Pred. No. 0.044;
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Score 43; DB 2;
Pred. No. 0.058;
Mismatches
                                                                                         DB 2; Length 71;
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              Indels
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         Gaps
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RESULT
$24215
Ig kappa chain - mouse (fragment)
C; Species: Mus musculus (house mouse)
C; Date: 19-Feb-1994 #sequence_revision 10-Nov-19;
C; Accession: $24215
R; Shimizu, T.; Iwasato, T.; Yamagishi, H.
J. Exp. Med. 173, 1065-1072, 1991
A; Title: Deletions of immunoglobulin C(kappa) re;
A; Reference number: $24214; MUID:91217618
A; Accession: $24215
A; Accession: $24215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT
34102
Ig kappa chain V region - human
C;Species: Homo sapiens (man)
C;Species: 13-Jan-1955 #sequence_revision 13-Ja
C;Accession: S34102
R;Wagner, S.D.; Luzzatto, L.
Eur. J. Immunol. 23, 391-397, 1993
A;Title: V-Kappa gene segments rearranged in
A;Reference number: S34076; MUID:93170387
A;Recession: S34102
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-78 < WAGS
A;Crossions S4106
PH1048
If light chain V region (clone 165.49) - mouse (fragment)
C:Species: Mus musculus (house mouse)
C:Date: 30-Sep-1993 *sequence_revision 30-Sep-1993 *text_(C:Accession: PH1048
R;Tillman, D.M.: JOU, N.T.: Hill, R.J.: Marion, T.N.
J. Exp. Med. 176, 761-779, 1992
A:Tille: Both 19M and 19G anti-DNA antibodies are the proc A:Reference number: PH0971; MUID:92381444
A:Accession: PH1048
A:Status: nucleic acid sequence not shown
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C;Superfamily: immunoglobulin V region; imm
C;Keywords: heterotetramer; immunoglobulin
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A; Residues: 1-79 <SHI>
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A;Cross-references: EMBL:X54832
C;Superfamily: immunoglobulin V region; immunoglobulin C;Keywords: heterotetramer; immunoglobulin F;7-Bl/Domain: immunoglobulin homology <IMM>
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C:Species: Homo sapiens (man)
C:Date: 19-Feb-1994 #sequence_revision 10-Nov-1995
C:Accession: S16834
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S16834
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C;Superfamily: immunoglobulin V region; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
E;7-81/Domain: immunoglobulin homology <IM
                                                                                  A;Status: preliminary; translation
A;Molecule type: mRNA
A;Residues: 1-86 <BLA>
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Eur. J. Immunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III
A;Reference number: S16823; MUID:91243737
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A;Reterence number: S34076; MUID:93170387
A;Accession: S34101
A;Status: preliminary
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A;Molecule type: mRNA
A;Residues: 1-86 < BLA>
A;Cross references: EMBL:X54824; NID:g33653; PIDN:CAA38593.1;
R;Wagner, S.D.; Luzzatto, L.
Eur. J. Immunol. 23, 391-397, 1993
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A;Residues: 1-81 <TIL)
A;Experimental source: B cell, strain [NZB x NZW]F1
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: immunoglobulin
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Pred. No. 0.067;
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334086

334086

If kappa chain V region - human
C:Species: Homo sapiens (man)
C:Date: 13-Jan-1995 *sequence_revision 13-Jan-1995 *text_change 21-Jan-2000
C:Accession: S34086
R:Wagner, S.D.; Luzzatto, L.
Eur. J. Immunol. 23, 391-397, 1993
A:Title: V-kappa gene segments rearranged in chronic lymphocytic leukemia are di
A:Reference number: S34076; MUID:93170387
A:Ccession: S34086
A:Status: preliminary
A:Molecule type: DNA
A:Residues: 1-86 <WAG>
A:Cross-references: EMBL:X67169
C:Superfamily: immunoglobulin V region; immunoglobulin homology
C:Keywords: heterotetramer; immunoglobulin homology
F:9-83/Domain: immunoglobulin homology <IMM>
RESULT 10

$16840

19 kappa chain V region - human (fragment)

C:Species: Homo sapiens (man)

C:Date: 19-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
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C;Species: Homo sapiens (man)
C;Date: 19-Feb-1994 *sequence_revision 10-Nov-1995 *text_change 21-Jan-2000
C;Accession: S16836
R;Blaison, G; Kuntz, J.L.; Pasquali, J.L.
Eur. J. Immunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III variable regions of polyclonal r)
A;Reference number: S16823; MUID: 91243737
A;Accession: S16836
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C;Superfamily: immunoglobulin V region; immunoglobulin C;Keywords: heterotetramer; immunoglobulin F;7-81/Domain: immunoglobulin homology <IMM>
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A; Residues: 1-86 <BLA>
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Ig kappa chain V region - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 19-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
C;Accession: $16837
R;Blaison, G; Kuntz, J.L.; Pasquali, J.L.
Eur. J. Immunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III variable regions of polyclonal rheumatoid A;Reference number: $16823; MUID:91243737
A;Accession: $1687
A;Accession: $1687
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-86 CBLA>
A;Residues: 1-86 CBLA>
A;Residues: Data: MEASA
                                                                                                                                                                                                     R;Blaison, G.; Kuntz, J.L.; Pasquali, J.L.
Eur. J. Immunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III variable regions of polyclonal rheumatoid
A;Reference number: S16823; MUID:91243737
A;Accession: S16833
                                                                                                                                                                                                                                                                                                                                C;Species: Homo sapiens (man)
C;Date: 19-Feb-1994 *sequence_revision 10-Nov-1995
C;Accession: S16833; S16838
                                                               A; Experimental source: clone bkv17
A; Accession: S16838
                                                                                                                A:Molecule type: mRNA
A:Residues: 1-86 <BLA>
A:Cross-references: EMBL:X54831
                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT
S16833
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C:Superfamily: immunoglobulin V region; immunoglobulin homology
C:Keywords: heterotetramer; immunoglobulin
F:7-81/Domain: immunoglobulin homology <IMM>
A; Molecule type:
A; Residues: 1-86
                                          A; Status: translation not shown
                                                                                                                                                                                       A; Status: translation not shown
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C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;7-81/Domain: immunoglobulin homology <TMM>
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A; Residues: 1-86 <BLA>
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R;Blaison, G.; Kuntz, J.L.; Pasquali, J.L.
Eur. J. Immunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III variable regions of polyclonal rheumatoid A;Reference number: S16823; MUID:91243737
A;Accession: S16840
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Matches 9; Conserv
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Pred. No. 0.071;
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RESULT 14

S16824

Ig kappa chain V region - human (fragment)

C;Specles: Homo saplens (man)
C;Specles: Homo saplens (man)
C;Date: 19:Feb-1994 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
C;Accession: S16824
R;Blaison, G; Kuntz, J.L.; Pasquali, J.L.
Eur, J. Immunol. 21, 1221-1227, 1991
A;Title: Molecular analysis of V(kappa)III variable regions of polyclonal rheumatoid fac A;Reference number: S16823; MUID:91243737
A;Status: preliminary; translation not shown
A;Molecule type: mRNA
A;Cross-references: EMBL:X54822
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin homology <IMM>
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C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Accession: $16830
R; Blaison, G.; Kuntz, J.L.; Pasquall, J.L.
Eur., J. Immunol. 21, 1221-1227, 1991
A; Title: Molecular analysis of V(kappa)III variable regions of polyclonal rheuma A; Reference number: $16823; MUID:91243737
A; Accession: $16830
A; Status: preliminary: translation not shown
A; Residues: 1-186 < BLAD: A; Rocession: $1680 < BNAD: A; Rocession: $1680 < BNAD: A; Rocession: $1680 < BNAD: A; Cross-references: EMBL: X4828
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 7-81/Domain: immunoglobulin homology < IMMD>
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A;Experimental source: clone sikv14
A;Experimental source: clone sikv14
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
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  TDFTLTISS 68
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ilarity 100.0%;
Conservative 0
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Pred. No.
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Pred. No. 0.071;
Mismatches 0
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Pred. No. 0.071;
; Mismatches
                                                                                Mismatches
                                                                                                     0.071;
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Ig kappa chain V region - human (fragment)
C;Speciles: Homo sapiens (man)
C;Speciles: Homo sapiens (man)
C;Date: 19-Feb-194 **sequence_revision 10-Nov-1995 **text_change 21-Jan-2000
C;Accession: S16829
C;Accession: S16829
C;Accession: S16829
C;Accession: S16823; MID:9127, 1991
A;Title: Molecular analysis of V(kappa)III variable regions of polyclonal rh
A;Reference number: S16823; MID:91243737
A;Accession: S16829
A;Status: preliminar; translation not shown
A;Status: preliminar; translation not shown
                                                                                                                                                                            Query Match
Best Local Similarity
Conserver
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C;Superfamily: immunoglobulin V region; immunoglobulin C;Keywords: heterotetramer; immunoglobulin E;7-81/Domain: immunoglobulin homology <IMM>
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S16829
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A; Residues: 1-86 <BLA>
Search completed: July 15, 2002, 13:01:02 Job time: 468 sec
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Perfect score:
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                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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ROCCOSETE
                                                                  KV1H_HUMAN STANDARD; PRT; 108 AA. P01600; 21-JUL-1986 (Rel. 01, Created) 21-JUL-1999 (Rel. 01, Last sequence update) 15-JUL-1999 (Rel. 38, Last annotation update) 1g kappa chain V-I region Hau.
                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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SMART; SMO0406; IGv; 1.
Immunoglobulin V region; Glycoprotein.
CARBOHYD 28 N-LINKED (GLCNAC.
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21-JUL-1986 (Rel. 01, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
15 kappa chain V-I region CAR.
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HSSP; P80362; 1WTL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE-75075135; PubMed-4216454;
Milstein C.P., Deverson E.V.;
"Primary structure of kappa light chain from a human myeloma
protein.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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InterPro; IPR003596; Ig_v
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WISCELLANEOUS: THE C REGION OF THIS CHAIN HAS THE INV (1,2)
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21-3UL-1986 (Rel. 01, Last sequence update)
15-3UL-1999 (Rel. 38, Last annotation update)
15-suppa chain V-V region UPC 61.
Mus musculus (Mouse).
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Watanabe S., Hilschmann N.;
"The primary structure of a monoclonal kappa-type immunoglobulin L-
"The primary structure of a monoclonal kappa-type immunoglobulin L-
                                                                                                                                                                                          InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; Ig; 1.
SMART; SM00406; IGV; 1.
                                                                                                                                                                                                                                                                                         PIR; A01929; KVMS61.
                                                                                                                                                                                                                                                                                                                                                 Vrana M., Rudikoff S., Potter M.;
"The structural basis of a hapten-inhibitable kappa-chain idiotype.";
J. Immunol. 122:1905-1910(1979).
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                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
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Mammalia; Eutheria;
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SMART; SM00406; IGV;
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1 23 FRAMEWORK-1.

24 34 COMPLEMENTARITY-D

35 49 FRAMEWORK-2.

50 56 COMPLEMENTARITY-D

57 88 FRAMEWORK-3.

88 97 COMPLEMENTARITY-D

98 107 FRAMEWORK-4.

23 88 BY SIMILARITY.
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Rodentia;
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COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
COMPLEMENTARITY-DETERMINING-3.
FRAMEWORK-4.
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COMPLEMENTARITY-DETERMINING-3
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COMPLEMENTARITY-DETERMINING-2
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KV5R_MOUSE
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      ROCCOSTTAN
                                                                                                                                                                                                             KV5S_MOUSE
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                                                                                                                                                                                                                                  RESULT
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Best Loc
Matches
Mis musculus (Mouse).

Eukaryota; Metazoa; Chordata; C.
Mammalia; Eutheria; Rodentia; S.
NCBI TaxID-10090;
[1]
                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
                                                                                            P01652;
21-JUL-1986 (Rel. 01
21-JUL-1986 (Rel. 01
15-JUL-1999 (Rel. 3E
1g kappa chain V-V I
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21-JUL-1986 (Rel.
21-JUL-1986 (Rel.
15-JUL-1999 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DOMAIN
DISULFID
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DOMAIN
DOMAIN
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J. Immunol. 122:1905-1910(1979).
-i- MISCELLANEOUS: THIS CHAIN WAS ISOLATED FROM MYELOMA PROTEINS THAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus (Mouse).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
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SMART; SM00406; IGV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PIR; B92808; KVMS09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=79195288; PubMed=109517; Vrana M., Rudikoff S., Potter M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HSSP; P01607;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KV5R_MOUSE
                                                                                                                                                                                             KV5S_MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NON_TER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR003006;
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9; Conser
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ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                           108 AA;
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                                                                                                                                                                                             STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01,
38,
                                                                                                01, Last seguence up 38, Last annotation V region J606.
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annotation update)
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COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPLEMENTARITY-DETERMINING-3 FRAMEWORK-4.
BY SIMILARITY.
                                          Craniata; Ver
Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                               Score 43; DB 1;
Pred. No. 0.015;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FAE4DA36076F2AFE CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           35C116BD60F79310 CRC64;
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                                                            Vertebrata;
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                                              Muridae;
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                                                              Euteleostomi;
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RESULT
KV5T_M
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Best Local (
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SMART; SM00406; IGV; 1.

Immunoglobulin V region.

DOMAIN 1 23

DOMAIN 35 49

DOMAIN 50 56

DOMAIN 57 88

DOMAIN 98 108

DISULFID 23 88

NON_TER 108 3. 1180
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21-JUL-1986
15-JUL-1999
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NON_TER
SEQUENCE
                  DOMAIN
DISULFID
                                                                   DOMAIN
                                                                                                       DOMAIN
DOMAIN
                                                                                                                                         DOMAIN
                                                                                                                                                                                                                                                                                                                                                MEDLINE-82099361; PubMed-6798111;
Johnson N., Slankard J., Paul L.,
"The complete V domain amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KV5T_MOUSE
P01653;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PIR; A92811; KVMS06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             binding proteins.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE-82099361; PubMed-6798111;
Johnson N., Slankard J., Paul L.,
"The complete V domain amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HSSP; P01607; 1REI
                                                                                                                                                                     InterPro; IPR003596; Pfam; PF00047; ig; 1 SMART; SM00406; IGV;
                                                                                                                                                                                                                                                               PIR; B92811
                                                                                                                                                                                                                                                                                                                               binding proteins."
                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                HSSP; P80362; 1WTL
                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID-10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ig kappa chain V-V region W3082
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro;
                                                      DOMAIN
                                                                                                                                                                                                                              InterPro; IPR003006;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    69 TOFTLTISS 77
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                                                                                                                                                                                                                                                                                                                  Immunol.
                                                                                                                                                                                                                                                                             Immunol. 128:302-307(1982).

MISCELLANEOUS: THIS CHAIN WAS :
BIND BETA(2-1)-FRUCTOFURANOSYL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MISCELLANEOUS: THIS CHAIN WAS BIND BETA(2-1)-FRUCTOFURANOSYL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      l Similarity
9; Conserv
                                                                                                                                                          lobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IPR003006;
IPR003596;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (Rel.
                                                                                                                                                                                                                                                               , KVMS82.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STANDARD;
region.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chordata;
Rodentia;
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                                                                                                                                                                                                           Ig_v.
                                                                                                                                                                                                                              Ig_MHC.
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                COMPLEMENTARITY-DETERMINING-1.
FRAMEMORK-2.
COMPLEMENTARITY-DETERMINING-2.
FRAMEMORK-3.
COMPLEMENTARITY-DETERMINING-3.
FRAMEMORK-4.
BY SIMILARITY.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPLEMENTARITY-DETERMINING-3 FRAMEWORK-4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FRAMEWORK-3
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COMPLEMENTARITY - DETERMINING - 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FRAMEWORK-1.

COMPLEMENTARITY-DETERMINING-1.
                                                                                                                                           FRAMEWORK-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Craniata; Ver
Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
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                                                                                                                                                                                                                                                                                                                                                Hood L.;
sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vertebrata;
thi; Muridae;
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RESULT 8
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SEQUENCE
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21-JUL-1986
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Primary structure of the variable part of an amyloidogenic Bence-Jones Protein (Mev). An unusual insertion in the third hypervariable region of a human kappa-immunoglobulin light chain.";

Hoppe-Seyler's Z. Physiol. Chem. 363:1347-1358(1982).
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SMART; SM00406; IG
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
Eukaryota;
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  Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
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BY SIMILARITY.
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Salomon A.;
Submitted (AUG-1996) to the SWISS-PROT data bank.
-1- MISCELLANEOUS: THE C REGION OF THIS CHAIN HAS
-1- MISCELLANEOUS: THIS IS A BENCE-JONES PROTEIN.
PTR; A01903: K4HULN.
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01-0C7-1996 (Rel. 34, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Ig kappa chain V-IV region Len.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Verte
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P01625;
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"The primary structure of a monoclonic immunoglobulin-L-chain of subgroup IV of the Rappa type (Bence-Jones protein Len).";
subgroup IV of the Rappa type (Bence-Jones protein Len).";
hoppe-Seyler's Z. Physiol. Chem. 356:507-557(1975).
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Mammalia; Eutheria;
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the European Bioinformatics Institute. The
use by non-profit institutions as long
modified and this statement is not removed.
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MEDLINE-81241357; PubMed-6788890;
Kwan S.-P., Rudikoff S., Seidman J.G.,
"Nucleic acid and protein sequences of
                                                                                          REVISION TO 9.
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NCBI_TaxID~10090;
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COMPLEMENTARITY-DETERMINING-1
                                                                                                                                                                                                                                              Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                            Craniata; Vertebrata; Euteleostomi;
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Pred. No.
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13-AUG-1987 (Rel. 05, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation updat
Ig kappa chain V-III region VG precursor (#Uman)
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                                                                                                                                                                                                                                                                                                                                                                  modified and this statement is not removed entitles requires a license agreement (See
                                                                                                                                                                                                                                                                                                             EMBL; X01668; -; NOT_ANNOTATED_CDS
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                                                                                                                                                                                                                                                                                                                                                                                                      the European Bioinformatics Institute. There are no rest
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Nucleic Acids Res. 12:9229-9236(1984).
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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SMART; SM00406; IGv; 1.
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COMPLEMENTARITY-DETERMINING-3
                                                                     FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
                                                                                                         IG KAPPA CHAIN V-III REGION VG FRAMEWORK-1.
COMPLEMENTARITY-DETERMINING-1.
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Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
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HSSP; P01789; 2MCP.
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Mammalia; Eutheria;
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13-AUG-1987 (Rel. 05, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Ig kappa chain V-III region VH precursor (Fra
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FRAMEWORK-1.
COMPLEMENTARITY-DETERMINING-1.
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21-JUL-1986 (Rel.
15-JUL-1999 (Rel.
                                                                                                                                                                                                                                                                                                                                                       Pfam; PF00047; ig; 1.
SMART; SM00406; IGV; 1.
Tmmunoqlobulin V region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; K01322; AAA58930.1; -.
EMBL; K01324; AAA58932.1; -.
EMBL; V00558; CAA23824.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bentley D.L., Rabbitts T.H.;
"Evolution of immunoglobulin V genes: evidence indicating
recently duplicated human V kappa sequences have diverged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Ig kappa chain V-I region HK101 precursor (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                              InterPro; IPR003006; Ig_MHC InterPro; IPR003596; Ig_v
                                                                                                                                                                                                                                                                                                                                                                                                                         HSSP; P01607; 1REI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
MEDLINE-83129397; PubMed-6402305;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                   A21056; A21056.
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9; Conser
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Last sequence update)
Last annotation update)
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COMPLEMENTARITY - DETERMINING - 2.
                                                                                                                                                                                                                                                                                                           FRAMEWORK-1.
COMPLEMENTARITY-DETERMINING-1.
                                                                                                                                                                                                                                                 COMPLEMENTARITY-DETERMINING-3.
BY SIMILARITY.
                                                                                                                                                                                                                                                                        FRAMEWORK-3
                                                                                                                                                                         Score 43;
Pred. No.
                                                                                                                                                                                                                                                                                                                                   IG KAPPA CHAIN V-I REGION HK101.
                                                   PRT;
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RESULT 14

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SEQUENCE
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                                                                                                            Homo sapiens (Human).
Eukaryota; Metazoa; C
Mammalia; Eutheria; F
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Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebr
             Klobeck H.G., Combristo G., Zachau H.G.; "Immunoglobulin genes of the Kappa light c lymphoid cell lines are closely related."; Nucleic Acids Res. 12:6995-7006(1984).
                                                            SEQUENCE FROM N.A. MEDLINE-85014148; Pubmed-6091049;
                                                                                                                                            13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
15-UUL-1999 (Rel. 38, Last annotation update)
Ig kappa chain V-I region Walker precursor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SIGNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                or send
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Subgroup IV of human immunoglobulin K light chains is encoded by single germline gene.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE-86041853; PubMed-2997712;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
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                                                                                                  NCBI_TaxID=9606;
                                                                                                                                                                                                  P04431;
                                                                                                                                                                                                              KV1W_HUMAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Klobeck H.G., Bornkamm
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9; Conserv
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Primates;
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Pred. No.
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COMPLEMENTARITY-DETERMINING-3
BY SIMILARITY.
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COMPLEMENTARITY-DETERMINING-2
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                                                                                                             Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                              PRT;
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RESULT 15
KV4B_HUMAN
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Best Local
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P06313;
01-JAN-1988
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                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
MEDLINE-86041853; PubMed-2997712;
Klobeck H.G., Bornkamm G.W., Comb
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SMART; SM00406; IGv; 1.
Immunoglobulin V region;
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                                                           EMBL; Z00022; CAA77317.1; PIR; A01904; K4HUJI.
                                                                                                                                                               use by non-profit institutions as long modified and this statement is not removed
                                                                                                                                                                                                                                                                                        single germline gene.
Nucleic Acids Res. 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ol.-JAN-1988 (Rel. 06, Created)
Ol.-JAN-1988 (Rel. 06, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation updat
Ig kappa chain V-IV region JI precursor.
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                                         HSSP; P01789; 1MCP
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COMPLEMENTARITY-DETERMINING-1.
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Pred. No. 0.0
Mismatches
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COMPLEMENTARITY-DETERMINING-2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     update)
                                                                                                                                                                                                                                                                                                                                                                            G., Mocikat R., Pohlenz H.D.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo
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InterPro;

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Search completed: July 15, 2002, 13:23:33 Job time: 1449 sec
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R SMART; SM00406; IGV; 1.

RIMULOGIODULIIN V region; S1,
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HAIN 21 43 FRA
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'IN 44 60 COMPL.
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83 114 FRAMENA.
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                                                                                            Query Match 100.0%;
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Matches 9; Conservative 0;
                                                                                                                                               Immunoglobulin V
SIGNAL 1
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DOMAIN 44
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SEQUENCE 133 J
                                                  1 TDFTLTISS 9
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95 TDFTLTISS 103
                                                                                                                                                  14632 MW; 5FB3953066744AF4 CRC64;
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COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
COMPLEMENTARITY-DETERMINING-3.
FRAMEWORK-4.
BY SIMILARITY.
                                                                                            ; Score 43; DB 1; Length 13; Pred. No. 0.018; 0; Mismatches 0; Indels
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1: sp_archea:*
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sp_bacteria:*
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Pred, No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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3 76.7	4 79.1	5 81.4	5 81.4	6 83.7	7 86.0	9 90.7	9 90.7	9 90.7	9 90.7	9 90.7	9 90.7	3 100.0	3 100.0	3 100.0	3 100.0	Query
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RESULT 2 Q9ERZ9 ID Q9ERZ9 PRELIMINARY; PRT; 107 AA.	TOPTLTISS 9	Ouery Match 100.0%; Score 43; DB 4; Length 107; Best Local Similarity 100.0%; Pred. No. 0.16; Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps	SQ SEQUENCE 107 AA; 11520 MW; 4BB43E9C5B577F16 CRC64;	NON_TER 107	DR EMBL; U96396; AAB68785.1;			RT "Molecular analysis of polyreactive monoclonal antibodies from		RX MEDLINE=98375893; PubMed=9712075;			Mammalia; Eutheria; Primates;	OS Homo Sapiens (Human). OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	DE VARIABLE REGION (FRAGMENT).		01-DEC-2001 (TrEMBLrel. 19, Last annotation update)	01-DEC-2001 (TrEMBLrel. 19,	01-DEC-2001 (TrEMBLrel.	ID Q96SA9 PRELIMINARY; PRT; IO/ AA.	6SA9	RESULT 1

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SMART; SM00406; IG; I
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Eukaryota; Metazoa; CF
Mammalia; Eutheria; Rc
NCBI_TaxID=10090;
                                                                                                                                                                                                                    SEQUENCE FROM N.A.
MEDLINE-98277139; PubMed-9614934;
MEDLINE-98277139; PubMed-9614934;
Van X., Liu B., Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                    01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE INMUNOGLOBULIN LIGHT CHAIN VARIABLE
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  InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
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Chen P., Deng J.B., Wang Z.L., Han H., Yao L.B.,
"Construction and sequencing of the single-chain
human TNF-alpha specific monoclonal antibody.";
T. 4 Chun i Ta Hsueh Hsueh Pao 19:373-376(1998).
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01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last senocation update)
ANTI HUMAN TNF-ALPHA LIGHT CHAIN VARIABLE REGION (FRAGMENT).
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Chen P., Deng J.B., Wang Z.L., Han H., Su C.Z.;

"Cloning and sequencing of the light chain fragment genes of an anti-hTNF-a monoclonal antibody.";

Cell. Mol. Immunol. 12:21-26(1996).
                                                                                                                                                                                                                                                                                                               NCBI_TaxID=9606;
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                                                                                                                                                                            Myosin-reactive autoantibodies
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Sciurognathi; Muridae;
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Wu X., Liu B.,
Young D.C.;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
                                                                                                                                                                                                                                                                    Q9UL81;
                                  SEQUENCE FROM N.A. MEDLINE-98277139; PubMed=9614934;
                                                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
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MEDLINE-98277139; PubMed-9614934;
MEDLINE-98277139; PubMed-9614934;
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Mammalia; Eutheria; Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBLrel. 13, Last sequence update)
EMBLrel. 19, Last annotation update)
IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
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11738 MW;
                    der Merwe
                                                                                                            Chordata;
Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11633 MW;
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                  P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 43;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          87:184-192(1998)
                                                                                                            Craniata; Vo
Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 43; DB
Pred. No. 0.1
D; Mismatches
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Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                       PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                Kalis N.N.,
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                                                                                                                                                                                                                                                                                       107
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                                                                                                                             Vertebrata; Euteleostom1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB
                                                                                                            Hominidae;
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RESULT
Q9UL79
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Best Local Similarity 88.
Matches 8; Conservative
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Best Local
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Q9UL79; PRELIMINARY;
Q9UL79; O1-MAY-2000 (TrEMBLrel. 13, C
Q1-MAY-2000 (TrEMBLrel. 13, L
Q1-DEC-2001 (TrEMBLrel. 19, L
MYOSIN-REACTIVE IMMUNOGLOBULI
(FRAGMENT).
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EMBL;
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NON_TER
SEQUENCE
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01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
MEDLINE-98277139; PubMed-9614934;
Wu X., Liu B., Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          InterPro; IPR003006; Ig_wHC.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
NON_TER 1 1
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HSSP; P80362; 1WTL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               fetus.";
Clin. Immunol. Immunopathol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Young D.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
Mammalia; Eutheria;
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; Ig; 1.
SMART; SM00406; IGV; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Myosin-reactive autoantibodies in rheumatic
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                                                                                                                                                                                                                                   TEFTLTISS
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8; Conser
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108 AA;
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107 AA;
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ilarity 100.0%;
Conservative
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                  EMBLrel. 13, Created)
EMBLrel. 13, Last sequence update)
EMBLrel. 19, Last annotation update)
IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
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; 11501 MW;
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11834 MW;
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Primates;
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                                                                                                                                                                                                                                                                                                                    Score 39; DB
Pred. No. 1.1;
1; Mismatches
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                            9F9C5A92EBA96EEA CRC64;
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1.1;
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                    REGION
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RESULT
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Best Local S
Matches 8
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Best Local S
Matches 8
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EMBL; AF035028; AAD56264.1; -.
HSSP; P80362; IMTL.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; Ig; 1.
SMART; SM00406; IGV; 1.
NON_TER
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SEQUENCE
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SEQUENCE FROM N.A.
MEDLINE-98277139; PubMed-9614934;
MEDLINE-98277139; PubMed-9614934;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGV; 1.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Euthería; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                 fetus."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Young D.C.;
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Young D.C.;
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70
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TDFTLTIS
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8; Conserv
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109 AA;
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                                                                                        Conservative
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ilarity 100.0%;
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11928 MW;
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11787
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                                                                                                            Score 39; DB 4;
Pred. No. 1.1;
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Pred. No.
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                                                                                                                                                                                                       243325F72C7DAC83 CRC64;
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Best Local Similarity

Matches 8; Conserv
    NON_TER
NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                    (FRAGMENT).

Homo sapiens (Human).

""karyota; Metazoa; Chordata; (""karyota; Metazoa; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NON_TER
SEQUENCE
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SEQUENCE FROM N.A.
SEQUENCE 98277139; PubMed=9614934;
MEDLINE=98277139; PubMed=9614934;
                                                                                                            InterPro;
InterPro;
                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
MEDLINE=98277139; Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   O9UL78;
01.MAY-2000 (TrEMBLrel. 13, Created)
01.MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
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09UL85;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN KAPPA CHAIN VARIABLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=9606;
                                                                      SMART;
                                                                                          Pfam;
                                                                                                                                                     EMBL; AF035036; AAL HSSP; P80362; 1WTL.
                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=9606
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                                                                                                                                                                                                                tetus.
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%; SM00406; IGV; 1.
                                                                                                                                                                                                                                                              D.C.;
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                                                                 PF00047; 1g; 1.; SM00406; IGV;
                                                                                                                                                                          Immunol. Immunopathol. AF035036; AAD56272.1;
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                                                                                   IPR003006; Ig_MHC
IPR003596; Ig_v.
0047; ig; 1.
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Primates;
      11646 MW;
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Pred. No.
                                                                                                                                                                                              87:184-192(1998)
                                                                                                                                                                                                                                                                                                                                                                                    Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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      5F675C52EC7EE197 CRC64;
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Best Local
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Q9JI.74;
Q9JI.74;
Q1-OCT-2000 (Tremblrel. 15, C:
Q1-OCT-2001 (Tremblrel. 19, L:
Q1-DEC-2001 (Tremblrel. 19, L:
ANTI-MYOSIN IMMUNOGLOBULIN LI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN-SF370 / ATCC 700294 / SEROTYPE M1;

MEDLING-21192684; PubMed-11296295;

Ferretti JJ / McShan W.M., Addic D.J., Savic G. Primeaux C., Sezate S., Suvorov A.N., Kenton S., Lai H.S., Qian Y., Jia H.G., Najar F.Z., Ren O., Zhu H., Song L., Whi Yuan X., Clifton S.W., Roe B.A., McLaughlin R.;

"Complete genome sequence of an M1 strain of Streptococcus Proc. Natl. Acad. Sci. U.S.A. 98:4658-4663(2001).

EMBL; AE006628; ARA34738.1; -.

InterPro; IPRO00559; FTHFS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PUTATIVE FORMATE-TETRAHYDROFOLATE LIGASE (EC 6.3.4. FHS.2 OR SPY2085.
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Q99XR2;
  with cardiac myosin.";
Infect. Immun. 68:5803-5808(2000).
EMBL; AF206032; AAF69330.1; ...
HSSP; P80362; IMTL.
                                                                                  MEDLINE-20448942: PubMed-10992488; Malkiel S., Liao L., Cunningham M.W., Diamond B.; "T-cell-dependent antibody response to the dominant streptococcal polysaccharide, N-acetyl-glucosamine, streptococcal polysaccharide, N-acetyl-glucosamine,
                                                                                                                                                                              SEQUENCE FROM N.A. STRAIN-BALB/C;
                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pfam; PF01268; FTHFS; 1.
PROSITE; PS00722; FTHFS_2; 1.
Ligase; Complete proteome.
SEQUENCE 557 AA; 59053 MW;
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Bacteria; Firmicutes; E
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                                                                                                                                                                                                                                                                                                            Mus musculus (Mouse).
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nilarity 100.0%;
Conservative 0
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Rodentia;
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77.8%;
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LIGHT CHAIN VARIABLE REG
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Pred. No.
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT;
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OSSPEE
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   PubMed-11679669;
Glaser P., Frangeul L.,
Baquero F., Berche P.,
                                                                                               SEQUENCE FROM N.A.
STRAIN-CLIP 11262 / SEROVAR 6A;
                                                                                                                                                                                                                                 Bacteria; Firmicutes; B
Bacillus/Staphylococcus
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MEDLINE-21361171; PubMed-11468171;

MEDLINE-21361171; PubMed-11468171;

Commenzo R.L., Zhang Y., Martinez C., Osman K., Herrera G.A.;

"The tropism of organ involvement in primary systemic amyloidosis:
contributions of Ig V(L) germ line gene use and clonal plasma cell
                                                                                                                                                                                                     NCBI_TaxID=1642;
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EMBL; AF361758; AAK5140
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Mammalia; Eutheria; Primates;
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
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Pred. No. 4.2;
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C., Amend A. Chakraborty
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RA Domann E., Dominguez-Bernal G., Duchaud E., Durant L., Dussurget O.,
RA Entian K.-D., Fsihi H., Garcia-del Portillo F., Garrido P.,
RA Gautier L., Goebel W., Gomez-Lopez N., Hain T., Hauf J., Jackson D.,
Jones L.-M., Kaerst U., Kreft J., Kuhn M., Kunst F., Kurapkat G.,
RA Madueno E., Maitournam A., Mata Vicente J., Ng E., Nedjari H.,
RA Mordsiak G., Novella S., de Pablos B., Perez-Diaz J.-C., Purcell R.,
RA Memmel B., Rose M., Schlueter T., Simoes N., Tierrez A.,
RA Vazquez-Boland J.-A., Voss H., Wehland J., Cossart P.;
"Comparative genomics of Listeria species.";
RI "Comparative genomics of Listeria species.";
RI Science 294:849-852(2001).
REMEL; ALS96171; CAC97510.1; -.
RKH Hypothetical protein: Complete proteome.
SGUENCE 1806 AA; 195670 MW; ABC808A65D84972F CRC64;
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K1 GLYCOPROTEIN (
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Viruses, dsDNA viruses, no RNA st
Gammaherpesvirinae; Rhadinovirus.
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                      completed: July 15,
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                                                                                                                                                                                                                                                 Score 34;
Pred. No.
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Perfect score:
Sequence:
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2000 Compugen
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AAY52745
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22	22	AAW52237	AAW52231	22	AAR88717	AAR98497	AAR98495	AAW01268		\mathbf{m}		AAW62803	AAW01285	AAR54318	AAG93599	AAY56683	AAY34316	AAR72060	AAR72059	AAR72058	AAR72065	AAR72064	AAR72063	AAR72062	AAR72061	AAB67510	σ	99	566	99	99	96	AAW62806
LD1/2-	LD1-6-1	LD2-20-	LD2-11-V	TD1/2	n antibody	2E3 light	2F2 light cha	VL region of HIV n	Anti-HIV gpl20 imm			sequen	gion of HI	IV qp120 in	_	nized 3Ğ9 liq	ntibody 2.	16 VK-1 I		VK req	434 VK-1 req	K.17 VK-1 L	K.19 VK-1 I	K.9 VK-1 L c	_	chain	o acid sequen	ial peptide f	peptide f	Partial peptide fr	tial peptide	RSV F	Amino acid sequenc

ALIGNMENTS

AAR87043

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AAR87043 standard; Peptide;

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Anti-human IL-5 recombinant antibody - useful for preventing or reducing eosinophilia and for treating certain allergic diseases.
                        WPI; 1996-058412/06
                                                (CLLT ) CELLTECH THERAPEUTICS LTD
                                                            17-JUN-1994;
                                                                         16-JUN-1995;
                                                                                      28-DEC-1995
                                                                                                  W09535375-A1
                                                                                                              Homo sapiens
                                   Bodmer
                                                            94GB-0012230
                                                                         95WO-GB01411.
                                   WW,
                                   Emtage JS
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Humanised antibody; interleukin-5; IL-5; recombinant antibody; antibody engineering; monoclonal antibody; MAb; 39D10; CDR; complementarity determining region; light chain; framework; eosinophilia; allergy; asthma.

Human group I light chain framework 3.

25-JUN-1996 AAR87043;

(first entry)

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RESULT AAX52745
ID AAX52745
ID AAX5
XX AAX5
XX AAX5
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XX BAX5
XX Hume
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KW Hume
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CC containing the variable region of the H chain of a mouse monoclonal Ab CC containing the wariable region of the H chain of a mouse monoclonal Ab CC recognising human tissue factor (hTF) and the constant region of the H CC chain of a human Ab. The variable region is one of six specified CC sequences (which are the H chain variable regions from mouse monoclonal CC Ab's ATR-2,3,4,5,7 or 8). Also described are chimeric Ab light (L) CC chains containing the variable region of the L chain of a mouse monoclonal Ab recognising human tissue factor (hTF) and the constant CC region of the L chain of a human Ab, the variable regions from mouse specified sequences (which are the L chain variable regions from mouse CC monoclonal Ab's ATR-2,3,4,5,7 or 8). The chimeric Ab's can be used for the treatment and prevention of thrombotic disease, especially of disseminated intravascular coagulation (DIC). The humanised antibody has the high'hff binding activity of the mouse monoclonal antibody but
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Humanised antibody recognizing human tissue factor, used of disseminated intravascular coagulation -
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ATR-2; ATR-3; ATR-4; ATR-5; ATR-7; ATR-8; thrombotic disease; DIC;
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Pred. No.
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Best Local S
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                                             express the A33 antigen. The method comprises administering an anticancer agent (I) conjugated to an immunoglobulin product (II) that binds specifically to A33 and contains one or more of 13 specified complementarity determining regions (CDRs), given in AAB98262 to AAB98274. (I) has cytostatic activity. The method can be used for treating colon and stomach cancers. (II), or the nucleic acid encoding it, can be used directly, in unconjugated form, for immunotherapy of cancer, and, when labeled, for detection or diagnosis of diseases associated with A33 expression. AAH22218 to AAH32254 and AAB98230 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              greatly reduced immunogenicity. AAZ33001 to AAZ33091 and Y527007 to AAY52767 represent sequences used in the exemplification of the present
                                                                                                                                                                                                                         Treating cancers, particularly of stomach and colon, that express A33 antigen by administering conjugate of anticancer agent with specific
                                                                                                                                                                                                                                                                                                               (SLOK )
(SCRI )
                                                                                                                                                                                                                                                                                                                                                                  22-OCT-1999;
04-APR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cytostatic; anticancer; colon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; rabbit; humanised; A33 antigen; anti-A33 antigen antibody; immunoglobulin; complementarity determining region; CDR; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB98286;
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                                                                                                                                                                                      Claim 16; Page 40; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200130393-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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Sequence
                                                                                                                                                  The present invention desembers the A33 antigen
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                                                                                                                                                                                                                                                                                         Barbas CF,
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                                  AAB98321 represent
                                                                                                                                                                                                                immunoglobulin product
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SLOAN KETTERING INST CANCER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
llarity 100.0%;
Conservative (
                                                                                                                                                               invention describes a method for treating cancers that
                                                                                                                                                                                                                                                                                                                                                                 99US-0425638.
2000US-0543004.
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Query Match Best Local Similarity

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Matches 9; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Treating cancers, particularly of stomach and colon, that express A33 antigen by administering conjugate of anticancer agent with specific
                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                               present invention.
                                                                                                                                                                                                                                                                                                             AAB98321 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a method for treating cancers that express the A33 antigen. The method comprises administering an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antigen by administering conjugate immunoglobulin product -
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04-APR-2000; 2000US-0543004.
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                                                    Chimeric antibody; humanised; humanisation; mammalian antibody; antigen; immunoreact; anti A33 antigen antibody; immunoglobulin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mammalian antibodies, which can be used for the treatment of a variety of diseases. The present sequence represents an A33 antigen binding immunoglobulin product VLFR3 peptide which is given in an example from
                                                                                                           A33 antigen binding immunoglobulin product VLFR3 peptide SEQ ID
                                                                                                                                                   08-AUG-2001
                                                                                                                                                                                           AAB97668;
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                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; ilarity 100.0%; Conservative
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Mismatches
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Preparing humanized rabbit antibodies that specifically immunoreact with a particular antigen using display technology for expressing libraries of antibody domains and fine tuning variable domain regions
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                                                                                     03-DEC-1996;
                                                                                                                                         03-DEC-1997;
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04-APR-2000; 2000US-0543004
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                        (ABGE-) ABGENIX INC
                                                                                                                                                                                                   11-JUN-1998
                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                   kappa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             libraries of antibody domains and fine tuning variable domain regions
                                                                                                                                                                                                                                                                                                                                                                                             Human; immunoglobulin; Ig; transgenic; non-human mammal;
Inactivated endogenous Ig locus; B-Cell development;
human heavy chain Ig locus; micro constant region; J-kappa gene; V-kappa;
kappa light chain Ig locus; kappa constant region; J-kappa gene; V-kappa;
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nes 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 TDFTLTISS 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                  9605-0759620
                                                                                                                                            97WO-US23091
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 43;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               display technology for expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 32
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AAW80981
ID AAW8
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Best Local Similarity
"hes 9; Conserv
   VALUE OF STREET 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modified genome also has an inserted human heavy chain Ig locus in germiline configuration, the human heavy chain Ig locus comprising a human micro constant region and regulatory and switch sequences, human J-H genes, human D-H genes, and human V-H genes and an inserted human kappa light chain Ig locus in germline configuration, the human kappa 11ght chain Ig locus comprising a human kappa constant region, J-kappa genes, and V-kappa genes, where the number of V-H and V-kappa genes inserted are selected to restore normal B-cell development in the mammal. The transgenic animals have a near complete human Ig locus, including both a be used for the production of human antibodies when exposed to particular antigens e.g. when exposed to human IL-B, EGFR or TWF- alpha contribution of human antibodies when exposed to particular in produce antibodies to IL-B, EGFR or TWF- alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Green L,
Mendez M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transgenic Xenomice, created using the method of the invention. The specification describes a transgenic non-human mammal which has genome modifications that comprise an inactivated endogenous immunoglobulin (
Humanised antibodies against epidermal growth factor receptor, EGF-r
                                                           WPI; 1999-034712/03.
                                                                                                                                                                                                                                      05-MAY-1997;
                                                                                                                                                                                                                                                                                               05-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                       12-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              transforming growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Variable kappa light region 012 encoded amino acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New transgenic non-human mammals - having an inactivated immunoglobulin locus and a near complete human immunoglobulin locus, used for production of human antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1998-333314/29
                                                                                                                                                                          (ABGE-) ABGENIX
                                                                                                                                                                                                                                                                                                                                                                                                                WO9850433-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; epidermal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 respectively.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 TDFTLTISS 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jakobovits
                                                                                                                    Jakobovits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                      9708-0851362
                                                                                                                                                                                                                                                                                               98WO-US09160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              growth factor receptor; tumour; EGF
wth factor alpha; TGF-alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 76
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Pred. No. 0.32;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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Best Local
AAM62793-822 represent fragments of human antibodies produced by transgenic xenomice, created using the method of the invention. specification describes a transgenic non-human mammal which has modifications that comprise an inactivated endogenous immunogloby
                                                                                                                                         Disclosure;
                                                                                                                                                                                      New transgenic non-human mammals - having immunoglobulin locus and a near complete bused for production of human antibodies
                                                                                                                                                                                                                                                                                                   WPI; 1998-333314/29
                                                                                                                                                                                                                                                                                                                                                                                Green L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           veterinary) to treat solid tumours. EGF-r is overexpressed on many human solid tumour types, and the fully human antibodies (i.e. comprising and inhibit both epidermal growth factor (EGF) and transforming growth factor alpha (TGF-alpha) binding to EGF-r (known to lead to cellular proliferation and tumour growth). They can prevent tumour cell growth and, in combination with an antineoplastic agent, may eradicate established tumours. The fully human antibodies can minimise the immunogenic and allergic responses intrinsic to previous mouse/rat or mouse/rat-derived antibodies.
                                                                                                                                                                                                                                                                                                                                                         Mendez
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9824893-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Amino acid sequence of a human antibody fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-SEP-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW62807 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      production;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  kappa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; immunoglobulin; Ig;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The variable kappa light region 012 encoded amino acid was used in the production of anti-epidermal growth factor receptor (EGF-r)-antibodies. The antibodies can be administered therapeutically to patients (human controlled)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inactivated endogenous Ig
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    heavy chain Ig locus; micro constant region; J-H; D-H; V-H light chain Ig locus; kappa constant region; J-kappa gene;
                                                                                                                                                                                                                                                                                                                                                       X
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3; Page 105; 143pp; English
                                                                                                                                                                                                                                                                                                                                                                                Jakobovits
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                                                                                                                                      Page
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ilarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                           Klapholz S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                locus; B-cell development;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       transgenic; non-human mammal;
                                                                                                                                      English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        82
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Pred. No.
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                                                                                                                                                                                                                immunoglobulin locus
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                                                                                                                                                                                                                                                                                                                                                                           æ
     immunoglobulin
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                         genome
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    V-kappa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local s
Matches s
      antibody (MAb) D. MAb D recognises Vibrio cholera neuraminidase-resistant epitope of ganglioside GD2, from a human melanoma cell line. MAbs A, B and C (see AAW14(84-89) are mentioned in the specification, but are not part of the claims. MAbs A and B recognise antigens 3 and 11 resp., of a permenant human tumour cell line. MAb C also recognises an epitope of a tumour-associated antigen. These antigens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 heavy; light chain; monoclonal antibody; antigen 3; marker; melanoma; permenant human tumour cell line; tumour-associated antigen; epitope; gastrointestinal tumour; pancreatic carcinoma; diagnostic; therapeutic; gastrointestinal tumour; pancreatic carcinoma; diagnostic; therapeutic;
                                                                                                  AAW14490-91 are the heavy and light chains (respectively) of monoclonal
                                                                                                                                  Disclosure; Page 14; 19pp; German.
                                                                                                                                                                gastrointestinal tumour marker
                                                                                                                                                                                                              N-PSDB; AAT63508
                                                                                                                                                                               Monoclonal antibody to tumour-associated
                                                                                                                                                                                                                                                          Auerbach B,
                                                                                                                                                                                                                                                                                      (BEHW ) BEHRINGWERKE
                                                                                                                                                                                                                                                                                                                     24-MAR-1989;
                                                                                                                                                                                                                                                                                                                                                  21-MAR-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Monoclonal antibody D VK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-JAN-1997
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9; Conserv
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                                                                                                                                                                                                                                                        Bosslet K,
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                                                                                                                                                                                                                                                                                                                   89DE-3909799
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                                                                                                                                                                                                                                                                                      ΑG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                                        Sedlacek
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 43;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AA
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                                                                                                                                                                                                                                                        Seemann
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                            tumour cell line.
MAD D (AAT36665-T36666) recognises a Vibrio cholera neuraminidase-
resistant epitope of ganglioside GD2, from a human melanoma cell
                                                                                                                                                                                                                                             therapeutic purposes.

MAbs A, B and D are mentioned in the specification, but are not part of the claims.

MAb A (AAT36661-T36662) recognises antigen 3 of permenant human tumour cell line.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-MAR-1989;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   marker; antigen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR99878 standard;
                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                    MAb C (AAT36659-T36660) is a monoclonal antibody that recognises an epitope of a tumour-associated antigen occurring at high concn. in the serum of patients with gastrointestinal tumours, e.g. pancreatic carcinoma, and is thus useful as a tumour marker for diagnostic or
                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 14; 19pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                               gastrointestinal tumour marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Monoclonal antibody to tumour-associated antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N-PSDB; AAT36666
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BEHW ) BEHRINGWERKE AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-MAR-1990;
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                                                                                                                                                                                                                               MAD B (AAT36663-T36664)
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nes 9; Conserv
                                Local Similarity
nes 9; Conserv
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                                                                                                                         84
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                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bosslet K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry
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                                                  100.0%;
                                                                                                                                                                                                                               recognises antigen 11 of permenant human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sedlacek
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                                                                                                                                                                                                                    CC light chain Ig locus in germline configuration, the human heavy chain tegions alight chain Ig locus of the chain Ig locus comprising a human because the constant region and regulatory and switch sequences, human J-H ce light chain Ig locus comprising a human CC denin Ig locus comprising a human V-H genes, and human V-H genes and an inserted human kappa CC light chain Ig locus in germline configuration, the human kappa light chain Ig locus comprising a human kappa constant region, J-kappa genes, cand V-kappa genes, where the number of V-H and V-kappa genes inserted are selected to restore normal B-cell development in the mammal. The CC transgenic animals have a near complete human Ig locus, including both a constant heavy chain locus and a human kappa light chain locus. They can comparticular antigens e.g. when exposed to human il-8, EGFR or TNF- alpha comparticular antigens e.g. when exposed to human il-8, EGFR or TNF- alpha compared to human il-8, EGFR or TNF- alpha
   Matches
                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW62793-822 represent fragments of human antibodies produced by transgenic Xenomice, created using the method of the invention. The specification describes a transgenic non-human mammal which has genome modifications that comprise an inactivated endogenous immunoglobulin (Ig) locus, so that the mammal does not display normal B-cell development. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New transgenic non-human mammals - having an inactivated immunoglobulin locus and a near complete human immunoglobulin locus, used for production of human antibodies
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Mendez M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human heavy chain Ig locus; micro constant region; J-H; D-H; V-H kappa light chain Ig locus; kappa constant region; J-kappa gene;
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                                                                                                                                            Sequence
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inactivated endogenous Ig locus; B-cell development;
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9; Conser
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   Conservative
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   0;
                               Score 43; DB 19;
Pred. No. 0.37;
      Mismatches
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V-kappa;
   Gaps
         0;
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RESULT 1
AAY56654
ID AAY5
                                                                                                            Query Match
Best Local Similarity
Watches 9; Conserv
                                                                                                                                                                                      RESULT 13
AAW59619
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                                                                                              Ωy
                                                                                                                                                                                                                    are reshaped human antibodies comprising a heavy chain selected from 19A, 19B, 19C or 19D (see AAM59615-18), which are based on the GL Dp8B sequence, and a light chain selected from 19A, 19B, 19C or 19D (see AAW59620-21). Such engineered antibodies are neutralising; they inhibit vitus growth in vitro and in vivo in animal models of RSV infection. Nucleic acids encoding the human MADs, recombinant plasmids (see AAV41427-33) and host cells (e.g. COS, CHO, myeloma)
                                                                                                                                                                                                                                                                                                        This is the heavy chain amino acid sequence of germline (GL) pk9 antibody. The invention relates to the construction and use of human MADS specific for the fusion (F) protein of respiratory syncytial virus (RSV) to passively treat, prevent or detect RSV infection. Huj9A, Huj9E, Huj9C and Huj9D MADS are claimed. These
         AAY56654 standard; protein; 88 AA
                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Fig 3; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                            Monoclonal antibodies reactive with Respiratory Syncytial Virus useful for detection, prevention and treatment of RSV infections
                                                                                                                                                                                                             are provided
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SMIK ) SMITHKLINE BEECHAM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Monoclonal antibody; human; Hu19; engineered antibody; RSV; respiratory syncytial virus; fusion protein; infection; complementarity determining region; CDR; therapy; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW59619;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW59619 standard; Protein; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W09819704-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anti-RSV F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Region
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                                                                                                                                                                                    88 AA;
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                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           96US-0030149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97WO-US19203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /label- CDR1
50..57
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                                                                                                                                  100.0%;
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                                                                                                                                  Score 43; DB 1
Pred. No. 0.38;
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                                                                                                                        Mismatches
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                                                                                                                                              DB 19;
                                                                                                                        0;
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                                                                                                                        Indels
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Query Match
Best Local Similarity
Watches 9; Conserve
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                                                                                                                                                                                                                                                                                                                                                        The invention provides an antibody (Ab) comprising donor CDRs (complementarity determining regions) derived from a non-human antigen-specific donor antibody, and an acceptor framework from a non-human primate. The Abs are prepared by grafting CDRs from a non-human antigen-specific donor antibody onto homologous Old World ape or monkey acceptor frameworks. The Abs have reduced immunogenicity and are better tolerated in humans (because of the close similarity between the human and primate proteins), but retain the full antigen-binding affinity of the donor
                                                                          Complementarity determining region; antibody; primate; immunogenicity; old World ape; Old World monkey; antigen-binding affinity.
                                 WO9955369-A1
                                                      Pan troglodytes
                                                                                                            Partial peptide fragment of chimpanzee Vkappa cDNA clone 46-6
                                                                                                                                        15-FEB-2000
                                                                                                                                                                 AAY56655
                                                                                                                                                                                      AAY56655 standard;
                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 22; Page 70-71; 123pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antibodies containing donor complementarity determining regions and non-human primate acceptor frameworks, having reduced immunogenicity humans -
                                                                                                                                                                                                                                                                                                                                                                                antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-023265/02.
N-PSDB; AAZ39322.
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Pred. No. 0.3
0; Mismatches
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                                                                                                                               Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 9; Conservative 0;
                                                                                                                                                                                                                                  The invention provides an antibody (Ab) comprising donor CDRs (complementarity determining regions) derived from a non-human antigenspecific donor antibody, and an acceptor framework from a non-human primate. The Abs are prepared by grafting CDRs from a non-human antigenspecific donor antibody onto homologous Old World ape or monkey acceptor frameworks. The Abs have reduced immunogenicity and are better tolerated in humans (because of the close similarity between the human and primate proteins), but retain the full antigen-binding affinity of the donor antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                   Antibodies containing donor complementarity determining regions and non-human primate acceptor frameworks, having reduced immunogenicity in humans -  \\
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-023265/02.
N-PSDB; AAZ39323.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Taylor AH;
                                                                                                                                                                                                                                                                                                                                                                                        Claim 22; Page 71; 123pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (SMIK ) SMITHKLINE BEECHAM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-APR-1999;
                                                                                                                                                                                                           Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99WO-US09131
                                                                                                                                    0;
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DΒ
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seq length: 2000000000
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1: /cgn2_6/ptodata/2/laa/5A_COMB.pep:*

2: /cgn2_6/ptodata/2/laa/5B_COMB.pep:*

3: /cgn2_6/ptodata/2/laa/6A_COMB.pep:*

4: /cgn2_6/ptodata/2/laa/6B_COMB.pep:*

5: /cgn2_6/ptodata/2/laa/PCTUS_COMB.pep:*

5: /cgn2_6/ptodata/2/laa/backfiles1.pep:*
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2:905 Million cell updates/sec
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2000
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                              PCT-US91-02942-42
US-08-59-147-40
US-08-59-147-061-13
US-09-347-061-13
US-09-347-061-13
US-09-343-004-92
US-09-543-004-92
US-09-543-004-99
PCT-US91-02942-9
US-08-765-179B-10
US-08-765-179B-10
US-08-765-179B-10
US-08-554-840-13
US-08-554-840-13
US-08-554-840-13
US-08-554-840-13
US-08-554-840-13
US-08-554-840-13
US-08-594-840-15
US-08-894-595-106
US-08-899-575-106
US-08-899-575-89
US-08-899-575-89
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Compugen Ltd
Sequence 42, Appl
Sequence 13, Appl
Sequence 21, Appl
Sequence 22, Appl
Sequence 94, Appl
Sequence 94, Appl
Sequence 94, Appl
Sequence 94, Appl
Sequence 97, Appl
Sequence 10, Appl
Sequence 11, Appl
Sequence 12, Appl
Sequence 13, Appl
Sequence 13, Appl
Sequence 15, Appl
Sequence 16, Appl
Sequence 17, Appl
Sequence 17, Appl
Sequence 18, Appl
Sequence 18, Appl
Sequence 106, App
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Sequence 41,
Sequence 41,
Sequence 92,
Sequence 92,
Sequence 94,
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100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
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US-08-300-386A-66	US-08-276-852-105	US-08-276-852-104	US-08-276-852-84	US-08-276-852-82	US-08-477-728-103	US-07-634-278-103	PCT-US95-08743-85	PCT-US95-08743-83	US-09-240-274-165	US-09-240-274-159	US-08-397-411-8	US-08-899-575-85	US-08-899-575-83	US-08-899-575-85	US-08-899-575-83	US-08-276-852-85	US-08-276-852-83
Sequence 66, Appl			Sequence 84, Appi						-	Sequence 159, App	_			-	•	`	83

ALIGNMENTS

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RESULT 1
PCT-US91-02942-42
                                                                                                                                                ; MOLECULE TYPE: protein PCT-US91-02942-42
                                                    Query Match
Best Local S
Matches 9
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                                                                                                                                                                                                                                     REGISTRATION NUMBER: 30,353
REFERENCE/DOCKET NUMBER: 1011.0586600
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 466-0800
TELEPHONE: (202) 466-0800
TELEPAX: (202) 833-8716
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHREACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNBER: PCT/US91/02942
FILING DATE: 19910429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: ROTHLEIN, ROBERT
APPLICANT: ADAIR, JOHN R
APPLICANT: ATHMAL, DILIEST S
TITLE OF INVENTION: HUMANIZED CDR-GRAFTED ICAM-1 ANTIBODY
NUMBER OF SEQUENCES: 102
                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: GB 9009549.8
APPLICATION NUMBER: GB 9009549.8
FILING DATE: 27-APR-1990
ATTORNEY_AGENT INFORMATION:
NAME: FOX, SAM L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
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ZIP: 200
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CITY: Washington
STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE:
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                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                      LENGTH:
1 TOFTLTISS 9
                                                      Similarity
9; Conserv
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                                                                          Score 43; DB 5; Pred. No. 0.058;
                                                        Mismatches
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STRANDEDNESS:
TOPOLOGY: line
MOLECULE TYPE: F
US-08-470-139-13
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Best Local Similarity
These 9; Conserve
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US-08-470-139-13
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Patent No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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                             SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/569,147
FILLING DATE: 25-March-1996
CLASSIFICATION: 536
ATTGRNEY/AGENT INFORMATION:
NAME: Trujillo, Doreen Yatko
REGISTATION NUMBER: 35,719
BREFERRATION NUMBER: 35,719
                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMOUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHIIN Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/470,139
FILING DATE: 06 JUNE-1995
                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
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IUMBER OF SEQUENCE...
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: IBM PC compatible
TOUTER: IBM PC compatible
TOUTER: PC-DOS/MS-DOS
TO1ease #1.0,
ELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                       STREET: One Libert:
CITY: Philadelphia
                                                                                                                                                                                                                                                          ZIP: 19103
                                                                                                                                                                                                                                                                                             STATE:
                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & ADDRESSEE: No. 6180377ris, LLP STREET: One Liberty Place - 46th Floor
                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                       COUNTRY:
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NAME: TRUJILLO, DOREEN YATKO
REGISTRATION NUMBER: 35,719
REGISTRATION NUMBER: CARP-0044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acids
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6180377
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100.0%; Pred. No.
tive 0; Mismatc
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         35,719
3ER: CARP-0047
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         SEQ ID NO 92
LENGTH: 32
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                            APPLICANT: Carlos F. APPLICANT: Lloyd J. Old TITLE OF INVENTION:
                                                                                                                                                                                          Sequence 92, Application US/09425638A Patent No. 6342587
                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                    CURRENT APPLICATION NUMBER: US/09/425,638A
CURRENT FILING DATE: 1999-10-22
NUMBER OF SEQ ID NOS: 129
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                                                                                                                   FILE REFERENCE:
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Matches 9; Conserv
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CURRENT FILING DATE: 1999-07-02
NUMBER OF SEQ ID NOS: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Bodmer, Mark
APPLICANT: Athwal, Diljeet Singh
APPLICANT: Emtage, John Spencer
TITLE OF INVENTION: Interleukin-
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc_feature
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TYPE: amino acid
STRANDEDNESS: single
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nes 9; Conserv
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linear
E:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (215) 568-3100
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                                                                                                                 LUD 5630
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                                                                                                                           A33 ANTIGEN SPECIFIC IMMUNOGLOBULIN PRODUCTS AND USES THE
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Pred. No. 0.075;
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Pred. No. 0.075;
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US-09-425-638A-92

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CURRENT APPLICATION NUMBER: US/09/543,004
CURRENT FILING DATE: 2000-04-04
PRIOR APPLICATION NUMBER: 09/425,638
PRIOR FILING DATE: 1999-10-22
NUMBER OF SEQ ID NOS: 129
LENGTH: 32
LENGTH: 32
TYPE: PRT
RESULT 8
US-09-543-004-94
· Sequence 94, Application US/09543004
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Best Local Similarity
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US-09-425-638A-94
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Patent No. 6346249
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LENGTH: 32
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                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Carlos F. Barbas III, Christoph Rader, Gerd Ritter, Sydney Welt and APPLICANT: Lloyd J. Old TITLE OF INVENTION: A33 ANTIGEN SPECIFIC IMMUNOGLOBULIN PRODUCTS AND USES THER
                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Carlos F. Barbas III, Christoph Rader, Gerd Ritter, Sydney Welt and APPLICANT: Lloyd J. Old
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                            FEATURE:
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Pred. No. 0.075;
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Pred. No. 0.075;
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Pred. No. 0.075;
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Query Match
Best Local Similarity
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; ORGANISM: Homo sapiens
; FEATURE:
US-09-543-004-94
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; MOLECULE TYPE: PCT-US91-02942-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 9, Application PC/TUS9102942
GENERAL INFORMATION:
APPLICANT: ROTHLEIN, ROBERT
APPLICANT: ADAIR, JOHN R
APPLICANT: ATHWAL, DILJEET S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Carlos F.
APPLICANT: Lloyd J. Old
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 94
LENGTH: 32
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                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9009549.8
FILING DATE: 27-APR-1990
ATTORNEY/AGENT INFORMATION:
NAME: FOX, SAM L
REGISTRATION NUMBER: 30,353
REGISTRATION NUMBER: 1011.0586
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/543,004
CURRENT FILING DATE: 2000-04-04
PRIOR APPLICATION NUMBER: 09/425,638
PRIOR FILING DATE: 1999-10-22
                                                                                                                TELEFAX: (202) 833-87 INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: ADAIR, JOHN R
APPLICANT: ATHWAL, DILJEET S
TITLE OF INVENTION: HUMANIZED CDR-GRAFTED ICAM-1 ANTIBODY
                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: PFILING DATE: 19910429
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Washington STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13 TDFTLTISS 21
                                        TOPOLOGY:
                                                                                                                                              TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
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                                                                        LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20036
                                                    AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Sterne, Kessler, Goldstein & Fox 1225 Connecticut Ave. NW Suite 300
                                                                        50 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
                                                                                                                              (202)
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                                                                                                                            (202) 466-0800
202) 833-8716
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Floppy disk
                  peptide
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                                                                                                                                                                                      1011.0586600
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                                                                                                                                                                                                                                                                                                                                                                                            Version #1.25
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RESULT 11
US-08-765-179B-14
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Best Local Similarity
Watches 9; Conservi
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                                                                                                                                                                              Sequence 14,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No. 5854027
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Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: DE P 4.
FILLING DATE: 15-UUL-1994
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 64 amino acids
                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                              tent No.
                                                                          APPLICANT: STEINBACHER, BOYIS
APPLICANT: STEINBACHER, Stefan
TITLE OF INVENTION: PROCESS FOR MODIFYING
TITLE OF INVENTION: OF ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/02626
FILING DATE: 06-UUL-1995
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/765,179B
FILING DATE: 14-JAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nikaido, Marmelstein, Murray & Oram LLP
STREET: 655 Fifteenth Street N.W. Suite 330
                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: STEIPE, BOTIS
APPLICANT: STEINBACHER, Stefan
TITLE OF INVENTION: PROCESS FOI
TITLE OF INVENTION: OF ANTIBOD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Washington STATE: D.C.
CITY: Washington
             STREET:
                                                                                                                                                                                                                                                            38 TDFTLTISS 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A. ZIP: 20005-5701
                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24
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                                                                                                                                                              4, Application US/08765179B
5854027
             E: Nikaido, Marmelstein, Murray & Oram LLP 655 Fifteenth Street N.W. Suite 330
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ilarity 100.0%;
Conservative (
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Pred. No. 0.16;
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                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                              THE STABILITY
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; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-765-179B-14
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Matches 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: BLACK, Amelia
APPLICANT: HANNA, Nabil
APPLICANT: PADLAN, Eduardo A.
APPLICANT: NEWMAN, Roland A.
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                  CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35,030
                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                              REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-6620
                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 2: CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 64 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/I
FILING DATE: 06-JUL-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                    TELEFAX:
                                                                                                                                                                APPLICATION NUMBER: US/08/554,840 FILING DATE: 07-NOV-1995
                                                                                                                                                                                                                                                                                                                   STATE: Virginia COUNTRY: United States ZIP: 22313-1404
                                                                                                                                                                                                                                                                                                                                                       STREET: P.O. Box
CITY: Alexandria
STATE: Virginia
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FILING DATE: 14-JAN-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: DE P 44 25 115.7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                         E: Burns, Doane, Swecker & Mathis P.O. Box 1404
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                    (703)
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                836-2021
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HUMANIZED ANTIBODIES TO HUMAN 9p39,
COMPOSITIONS CONTAINING AND THERAPEUTIC USE THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCT/EP95/02626
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Pred. No. 0.16;
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SEQUENCE CHARACTERISTICS:

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APPLICANT: Yang, Xiao-Dong
APPLICANT: Yang, Xiao-Dong
APPLICANT: Gallo, Michael
APPLICANT: Gallo, Michael
APPLICANT: Jia, Xiao-Chi
TITLE OF INVENTION: Human Monoclonal Antibodies to Epi
TITLE OF INVENTION: Growth Factor Receptor
FILE REFERENCE: Cell 4.20
CURRENT APPLICATION NUMBER: US/08/851,362D
CURRENT FILING DATE: 1997-05-05
NUMBER OF SEQ ID NOS: 49
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 21
LENGTH: 76
TWYPE: DET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; MOLECULE TYPE: protein US-08-554-840-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TYPE: PRT
; ORGANISM: human
US-08-851-362D-21
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US-08-554-840-12
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Best Local Similarity
Matches 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                   sequence 12, Application US/08554840 Patent No. 6001358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
COUNTRY: United States
ZIP: 22313-1404
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Burns, Doane, Swecker & Mathis
STREET: P.O. Box 1404
                                                                                                                                                                                           APPLICANT: PADLAN, Eduardo A.
APPLICANT: NEWMAN, ROland A.
TITLE OF INVENTION: HUMANIZED ANTIBODIES TO HUMAN 9D39,
TITLE OF INVENTION: COMPOSITIONS CONTAINING AND THERAPEUTIC USE THEREOF
                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                 APPLICANT:
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TOPOLOGY: li
                                                                                                  STREET: P.O. Box
CITY: Alexandria
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                    50 TDFTLTISS 58
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mes 9; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                    1 TDFTLTISS 9
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                                                                                                                                                                                                                                                                 HANNA, Nabil
                                                                                                                                                                                                                                                                                 BLACK, Amelia
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Pred. No. 0.19;
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Pred. No. 0.17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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US-08-554-840-12
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                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/554,840
FILING DATE: 07-NOV-1995
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35,030
REGISTRATION NUMBER: 012712-127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LEWGTH: 80 amino acids
TYPE: amino acid
                                                                                                   INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION: NAME: Teskin, Robin L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: BLACK,
APPLICANT: HANNA,
APPLICANT: PADLAN
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-6620
                                                                                                                                TELEPHONE: (703) 836-6620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
MOLECULE TYPE:
                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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FILING DATE: 07-NOV-1995
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Teskin, Robin L. REGISTRATION NUMBER: 35,030
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CITY: Alexandria
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                                                              80 amino acids
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                                                                                                                (703) 836-2021
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ALIGNMENTS

RESULT

trans-activating transcription regulator - human immunodeficiency virus type 1 (isola C;Species: human immunodeficiency virus type 1, HIV-1 A;Note: host Homo sapiens (man)

C;Date: 30-Sep-1991 #sequence_revision 30-Sep-1991 #text_change 16-Jul-1999

Hampe,

C;Accession: JQ0071 R;Spire, B; Sire, J.; Zachar, V.; Rey, F.; Barre-Sinoussi, F.; Galibert, F.; Hamp, Gene 81, 275-284, 1989 Gene 81, 275-284, 1989 A;Title: Nucleotide sequence of HIV1-NDK: a highly cytopathic strain of the human A;Reference number: JQ0065; MUID:90034200 A;Accession: JQ0071 R;Michael, N.L.; Chang, G.; d'ARCY, L.A.; Ehrenberg, P.K.; Mariani, R.; Busch, M.P.; J. Virol. 69, 4228-4236, 1995 A;Title: Defective accessory genes in a human immunodeficioncy virus type 1-infected A;Reference number: Z16654; MUID:95287475 A;Accession: T99384 A;Status: preliminary; translated from GB/EMBL/DDBJ A;Molecule type: DNA A;Residues: 1-71 MICO trans-activating transcription regulator - human immunodeficiency virus type 1 C;Species: human immunodeficiency virus type 1, HIV-1 C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 21-Ju1-2000 C;Accession: T09384 Query Match Best Local Similarity "~+~hes 9; Conserva C; Superfamily: C; Keywords: AII A;Gene: tat C;Superfamily: AIDS trans-activating transcription regulator C;Keywords: transcription A;Cross-references: EMBL:U24451; NID:g829440; PIDN:AAA79576.1; PID:g829444 C;Genetics: 밁 δô A; Gene: tat C; Genetics: A;Molecule type: DNA A;Residues: 1-86 <SPI> A;Cross_references: GB:M27323; NID:g328154; PIDN:AAA44866.1; PID:g328155 47 1 YGRKKRRQKKK 11 N YGRKKRRQRRK 57 ly: AIDS trans-activating transcription AIDS; immunodeficiency; transcription Conservative 89.7%; 81.8%; Score 52; DB 1 Pred. No. 0.15; Mismatches DB 1; 0; regulator Length 86; Indels 0 Gaps 0 (isola

lymphoid enhancer T-cell receptor a unknown protein, 6 T-cell receptor al

Query Match

84.5%;

Score 49;

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Length 71;

DnaJ-like protein helicase homolog ribosomal Na+/K+-exchanging hypothetical prote

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A; Gene: tat
A; Introns: 72/3
C; Superfamily: A
C; Keywords: AIDS
                                                              A25700
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A; Gross-references: GB:M13137; NID:g326460
A: Note: the GenBank entry ADRE3AA PID:g209908
C; Genetics:
A; Gene: tat
C; Superfamily: AIDS trans-activating transcrip
C; Keywords: transcription regulation
trans-activating transcription regulator - human immunodeficiency virus type C;Species: human immunodeficiency virus type 1, HIV-1 C;Date: 28-Sep-1987 *sequence_revision 28-Sep-1987 *text_change 02-Jul-1998 ''''
                                                                                                                                                                                                                                                                                                                                                                                                                               Gene 52, 71-82, 1987
A:Title: Molecular characterization of human A:Reference number: A26192; MUID:87248097
A:Accession: C26192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    trans-activating transcription regulator - human immunodeficiency virus C; Species: human immunodeficiency virus Zr-6 C; Date: 30-Sep-1987 *sequence_revision 30-Sep-1987 *text_change 16-Jul-1 C; Accession: C26192
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A:Title: Molecular cloning and primary nucleotide sequence (A:Reference number: A94136; MUID:87041461
A:Accession: B25523
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                                                                                                                                  В
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A; Residues: 1-86 <SRI>
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8; Conservative
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A; Cross-references: E
C; Genetics:
A; Gene: tat
A; Introns: 72/2
C; Superfamily: AIDS t
                                                                                                                          A;Cross-references: EMBL;Z11530; NID:g60192; PIDN:CAA77625.1; PID:g60196
R;Siderovski, D.P.; Matsuyama, T.; Frigerio, E.; Chui, S.; Min, X.; Erfle, H.; Sumner
Nucleic Acids Res. 20, 5311-5320, 1992
A;Title: Random mutagenesis of the human immunodeficiency virus type-1 trans-activato
A;Reference number: S26385; MJID:93065196
A;Accession: S26385
                                                                                           A; Molecule type: nucle
A; Residues: 1-86 <SID>
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A; Residues: 1-86 <CAR>
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R;Carlini, F
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A; Residues: 1-86 <THE>
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R:Theodore, T.; Buckler-White, A.J.
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C;Species: human immunodeficiency virus type 1, HI
C;Date: 15-Jul-1995 #sequence_revision 01-Sep-1995
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S54381
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A;Residues: 1-86 <SOD>
C;Superfamily: AIDS trans-activating transcription
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R;Sodroski, J.; Patarca, R.; Rosen, C.; Wong-Staal, F.; Haseltine, Science 229, 74-77, 1985
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A:Residues: 1-87 <ALI>
A:Cross-references: EMBL:K03456; NID:g60228; PIDN:CAA28015.1;
C:Genetics:
A:Introns: 72/2
C:Superfamily: AIDS trans-activating transcription regulator
                                                                                                                                                                                                                                                                                                                                                          R;Alizon, M.; Wain-Hobson, S.; Gluckman, J.C.; Sonigo, P. Cell 46, 63-74, 1986
A;Title: Genetic variability of the AIDS virus: Nucleotide A;Reference number: Z14389; MUID:86245056
A;Accession: T01665
     trans-activating
C; Species: human
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C;Species: human immunodeficiency virus type 1, HIV-1
C;Date: 19-Feb-1999 #sequence_revision 19-Feb-1999 #text_change
C;Accession: T01665
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Blochem. Blophys. Res. Commun. 235, 806-811, 1997
A;Title: A selection system to study protein-RNA
A; Reference number: JC5591; MUID:97350867
A; Accession: JC5591
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C:Species: human immunodeficiency virus type 1, HIV-1
C:Date: 23-Sep-1997 #sequence_revision 23-Sep-1997 #text_change 17-Mar-1999
C:Accession: JC5591
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8; Conser
transcription regulator - human immunodeficiency virus type 1 (isolate immunodeficiency virus type 1, HIV-1 \,
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Pred. No.
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Pred. No. 0.46;
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Pred. No. 0.45;
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tat protein - human immunodeficiency virus type 1 (strain JRFL) C. Species: human immunodeficiency virus type 1, HIV-1 C. Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change C. Accession: T09446 F. Pang, S.; Vinters, H.V.; Akashi, T.; O'Brien, W.A.; Chen, I.S. submitted to the EMBL Data Library, July 1996 A. Reference number: Z16673 A. Accession: T09446
C;Genetics:
A;Gene: tat
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C;Superfamily: AIDS trans-activating transcription regulator
C;Keywords: AIDS; immunodeficiency; transcription regulation
                                   A;Cross-references: EMBL:U63632; NID:g1465777; PID:g1465783
                                                           A; Residues:
                                                                                A; Molecule type: DNA
                                                                                                      A; Status: preliminary; translated
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A;Gene: .tat
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A; Residues: 1-101 <LIY>
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A;Title: Complete nucleotide sequence, genome organization, and biological properties.
A;Reference number: A44001; MUID:93021387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N:Alternate names: tat protein
C:Species: human immunodeficiency virus type 1, HIV-1
A:Note: host Homo sapiens (man)
C:Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 02-Jul-1998
C:Accession: E44001
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C;Superfamily: AIDS trans-activating transcription
C;Keywords: AIDS; immunodeficiency; transcription
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A; Residues: 1-95 <ARY>
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A;Title: Three novel genes of human T-lymphotropic virus type
A;Reference number: 894093; MUID:86177573
A;Accession: A04017
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6;Date: 04-Dec-1986 #sequence_revision 04-Dec-1986 #text_change 02-Jul-1998
6;Accession: A04017
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Best Local :
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Pred. No. 0.52;
3; Mismatches
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C;Accession: Data 7; Sulton 7; Sulton 7; Sulton 7; Sulton 6; Accession: Data 95; Sulton 7; Sulton 7; Sulton 7; Sulton 7; Sulton 8530-8536, 1992
A;Title: The human r cell transcription factor-1 gene. Structure, localization, and p A;Accession: D38095
A;Accession: D38095
A;Status: preliminary
A;Accession: D38095
A;Status: preliminary
A;Accession: Selson
A;Cross -references: G8:X63901
A;Cross -references: G8:X63901
A;Cross -references: Selson
A;Accession: Science
A;Acce
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C;Date: 19-May-1994 #sequence_revision 19-May-1994 #text_change 24-Sep-1999
C;Accession: D38095; S61877; S61800
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                                                                                                          T-cell-specific transcription factor 1 splice form C - human NyAlternate names: transcription factor TCF-IC C:Species: Homo sapiens (man) C:Bete: 19-May-1994 *sequence_revision 19-May-1994 *text_change 24-Sep-1999 C:Accession: C:B095; S13449; S65007
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C; Superfamily: AIDS
R:van de Wetering, H.; Oosterwegel, M.; Holstege, J. Biol. Chem. 267, 8530-5536, 1992
A;Title: The human T cell transcription factor-1 (A;Reference number: A38095; MUID:92235082
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nes 8; Conserv
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                                                                                          F.; Dooyes, D.; Suijkerbuijk,
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A;Cross-references: EMBL:X59871; NID:g36789; PIDN:CAA42528.1; PID:g36790 A;Note: the authors describe an additional C-terminal exon of splice form C, erences A38095 and S61796 R;Mayer, K; Wolff, E; Clevers, H.; Ballhausen, W.G. Biochim. Biophys. Acta 1263, 169-172, 1995 A;Title: The human high mobility group (HMG)-box transcription factor TCF-1: A;Reference number: S61796; MUID:95367594
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Title: Identification and cloning of TCF-1, a T lymphocyte-specific transcription A;Reference number: S13449; MUID:91114695
A;Accession: S13449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R;van de Wetering, M.; Oosterwegel, M.; Holstege, F.; J. Blol. Chem. 267, 8530-8536, 1992
A;Title: The human T cell transcription factor-1 gene. A;Reference number: A38095; MUID:92235082
A;Accession: A38095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T-cell-specific transcription factor 1 splice form A - human N;Alternate names; transcription factor TCF-1A C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Date: 19-May-1994 *sequence_revision 19-May-1994 *text_change 21-Jul-2000 C;Accession: A38095; S16645; S65005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Superfamily: unassigned HMG box proteins; HMG box homology C;Rupwords: alternative splicing; DNA binding; transcription F;151-226/Domain: HMG box homology <HMGl>
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A; Residues: 244-268 <MAY>
A; Note: only a part of the
C; Genetics:
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EMBO J. 10, 123-132, 1991
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A; Residues: 1-268 < VAN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Introns: 32/3; 68/1; 97/2; 137/2; 191/3; 227/3 244/1 C; Superfamily: unassigned HMG box proteins; HMG box homology
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A; Status: not compared with A; Molecule type: mRNA A; Residues: 244-269 <MAY>
                                                                                                                         A;Cross-references: EMBL:X59869; NID:g36785; PIDN:CAA42526.1; R;Mayer, K.; Wolff, E.; Clevers, H.; Ballhausen, W.G. Blochim. Blophys. Acta 1263, 169-172, 1995
Blochim. Blophys. Acta 1263, 169-172, 1995
A;Title: The human high mobility group (HMG)-box transcription
                                                                                                                                                                                                                               A; Molecule type: mRNA
A; Residues: 1-269 <WET>
                                                                                                                                                                                                                                                                                       A; Reference number: A; Accession: S16645
                                                                                                                                                                                                                                                                                                              A; Title: Identification and cloning of TCF-1, A; Reference number: S13449; MUID: 91114695
                                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: GB:X63901; NID:g36791; PIDN:CAB56795.1; PID:g6006565 A;Note: authors translated the codon GAT for residue 253 as Ser R;van de Wetering, M.; Oosterwegel, M.; Dooijes, D.; Clevers, H. EMBO J. 10, 123-132, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-269 < VAN>
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7; Conserv
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                                                                                                        MUID:95367594
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Pred. No. 10;
3; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HMG)-box transcription factor TCF-1: novel
                                                                                                                                    (HMG)-box transcription
                                                           translation
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                                                                                                                                       factor TCF-1:
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C:Gemetics:
A:Introne: 30/1; 60/1; 97/2; 131/3; 121/3; 227/3; 244/1
C:Seperismily: unassigned BMG low Proteins; BMG low homology
C:Reywords: alternative splicing: DMR binding: transcription factor
F:151-226/homain: BMG low homology chedi:
Query Match
74.1%; Score 43: DB 2: Length 269;
Bast Local Similarity 63.6%; Pred. No. 10;
WACHES 7: Conservative 3; Mismatches 1; Indels 0; Gaps 0;
QY 1 YOMKEROSKK 11
DB 277 YOKKERSEEK 2177
Search completed: July 15, 2002, 13:01:03
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     Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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MEDLIND-85111123; PubMed-2578615;

Rather L., Haseltine W., Patarca R., Livak K.J., Starcich B.R.,

Josephs S.F., Doran B.R., Rafalski J.A., Whitehorn E.A.,

Josephs S.F., Ivanoff L., Petteway S.R. Jr., Pearson M.L.,

Lautenerger J.A., Papas T.S., Chrayeb J., Chang N.T., Gallo R.
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P04612;
              MEDLINE-86177573; PubMed-3008154;
                        SEQUENCE FROM N.A.
                                              NCBI_TaxID=11679;
                                                      Human immunodeficiency virus type 1 (clone 12) (HIV-1).
Viruses; Retroid viruses; Retroviridae; Lentivirus.
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SUBUNIT: BINDS CYCLIN T1 (BY SIMILARITY).
SUBCELLULAR LOCATION: NUCLEAR, NUCLEOLAR.
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   STRAIN-ISOLATE HXB3;
MEDILINE-85228248; PubMed-2988795;
MEDILINE-85228248; PubMed-2988795;
Crowl R., Ganguly K., Gordon M., Conroy R., Schaber M.,
Shaw G.M., Wong-Staal F., Reddy E.P.;
"HTLY-III env gene products synthesized in E. coli are I antibodies present in the sera of AIDS patients.";
Cell 41:979-986(1985).

Cell 41:979-986(1985).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TAT_HV1B1
P04606;
13-AUG-1987
13-AUG-1987
16-OCT-2001
                                                                                                                                                                                                                                                             "Complete nucleotide sequence of the AIDS virus, HTLV-III.";
Nature 313:277-284(1985).
                                                                                                                                                                                                                                                                                                                                                             Ratner L., Haseltine W., Patarca R., Livak K.J., Starcich Josephs S.F., Doran E.R., Rafalski J.A., Whitehorn E.A., Baumeister K., Ivanoff L., Petteway S.R. Jr., Pearson M.L. Lautenberger J.A., Papas T.S., Ghrayeb J., Chang N.T., Gal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human immunodeficiency virus type 1 (BH10 isolate)
Human immunodeficiency virus type 1 (HXB3 isolate)
Viruses; Retroid viruses; Retroviridae; Lentivirus
NCBI_TaxID=11678, 11707;
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InterPro; IPR001831; HIV_Tat.
Pfam; PF005539; Tat; 1.
PRINTS; PR00055; HIVTATDOMAIN.
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SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
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FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
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TAT_HV1BR
P04610;
                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its most profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.

MEDLINE-85099333; PubMed-2981635;

Main-Hobson S., Sonigo P., Danos O., Cole S., Alizon

"Nucleotide sequence of the AIDS virus, LAV.";
                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A. (CLONE PN14-3).

Buckler C.E., Buckler-White A.J., Willey R.L., McCoy J.;

Submitted (JUN-1988) to the EMBL/GenBank/DDBJ databases.

-i- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human immunodeficiency virus type 1 (BRU isolate) (HIV-1). Viruses; Retroid viruses; Retroviridae; Lentivirus.
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16-OCT-2001
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Pfam; PF00539; Tat; 1.
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HIV; M14100; TAT$HXB3.
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                                                                                                                                                                                                                                                                                            SUBUNIT: BINDS CYCLIN T1
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(Rel. 40, Last annotation updat
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Pred. No. 0.074;
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P04608; O09778;
13-AUG-1987 (Re
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ratner L., Fisher A., Jagodzínski L.L., Mitsuya H., Liou R.-S., Gallo R.C., Wong-Staal F.;
Submitted (APR-1997) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LI
                                                                                                                                                                                                            use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See
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HIV; K03455; TAT$HXB2.
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the European Bioinformatics Institute. There are no rest
use by non-profit institutions as long as its content
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-!- SUBUNIT: BINDS CYCLIN T1 (BY SIMILARITY).
-!- SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
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                                            InterPro; IPR001831; HIV_Tat.
Pfam; PF00539; Tat; 1.
                                                                                                                                          EMBL; K03455; AAB50256.1;
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NCBI_TaxID=11706;
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
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HIV; M19921; TAT$NL43
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16-OCT-2001
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Pred. No. 0.07
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Pram;

Transcription

regulation; Activator; RNA-binding; Nuclear protein;

PRINTS; PR00055; HIVTATDOMAIN

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RESULT 8
TAT_HV12%
ID TAT_H
AC P1250
DT 01-OC
                                                                                                                                                                                                                                                                                                                                                                      TAT_HV1PV
ID TAT_H
AC P0460
DT 13-AU
DT
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Best Local Similarity
Whiches 8; Conserv:
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Best Local S
Matches 8
TAT_HV1Z2
P12506;
01-OCT-1989
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P04607;
13-AUG-1987
13-AUG-1987
16-OCT-2001
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR001831; HIV_Tat Pfam; PF00539; Tat; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HIV; K02083; TAT$PV22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; K02083; AAB59870.1; -.
EMBL; X01762; -; NOT_ANNOTATED_CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Capon D.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SBQUENCE FROM N.A.
MEDLINE-85111157; PubMed-2982104;
Muesing M.A., Smith D.H., Cabradilla C.D., Benton C.V.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Viruses; Retroid viruses;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TAT protein
                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                   Transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                               PRINTS; PR00055; HIVTATDOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Nucleic acid structure and expression of the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus type 1 (PV22 isolate) (HIV-1).
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8; Conserv
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(Rel. 05, Last sequence update)
(Rel. 40, Last annotation update)
     (Rel. 12, Created)
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Pred. No.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TAT_HV126
P04609;
13-AUG-1987
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Submitted (NOV-1988) to the HIV data bank.

-i- FUNCTION: TRANSCRIETIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human immunodeficiency virus type 1 (Z2/CDC-Z34 isolate) (HIV-1). Viruses; Retroid viruses; Retroviridae; Lentivirus.
                                                                          "Molecular characterization of human immunodeficiency virus from Zaire: nucleotide sequence analysis identifies conserved and variable domains in the envelope gene.";

Gene 52:71-82(1987).
-!- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND RACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
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                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
MEDLINE-87248097; PubMed-3036660;
Srinivasan A., Anand R., York D., Ranganathan
Schochetman G., Curran J., Kalyanaraman V.S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AIDS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Viruses; Retroid viruses; Retroviridae; Lentivirus NCBI_TaxID=11708;
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16-OCT-2001
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SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
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  SUBUNIT: BINDS CYCLIN T1 (BY SIMILARITY) SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR
                                                        PROMOTER.
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(Rel. 05, Last sequence update)
(Rel. 40, Last annotation update)
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72.7%;
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Luciw P.A.,
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RESULT 10
TAT_HYIMA
ID TAT_HYIMA
ID TAT_H
AC P0461
DT 13-AU
DT 13-AU
DT 16-OC
DE TAT P
OS Human
OC Virus
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P04613;
13-AUG-1987
13-AUG-1987
                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
MEDLINE-86245056; PubMed-2424612;
Alizon M., Wain-Hobson S., Montagnier L., Sonigo P.;
"Genetic variability of the AIDS virus: nucleotide sequence analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human immunodeficiency virus type 1 (MAL isolate) (HIV-1). Viruses; Retroid viruses; Retroviridae; Lentivirus. NCBI_TaxID=11697;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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      SEQUENCE
                                                                                        PRINTS;
                                                                                                                                                                                               EMBL; X04415; CAA28015.1; -.
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                                                         Transcription
                                                                                                                                               InterPro;
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nes 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            two isolates from African patients.";
11 46:63-74(1986).
- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE
TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROMOTER.
SUBUNIT: BINDS CYCLIN T1 (BY SIMILARITY).
SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
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                                                                                                                               rPro; IPR001831; HIV_Tat.
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                                                                                                                PF00539; Tat;
                                                                                        PR00055; HIVTATDOMAIN
      87
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(Transactivating regulatory protein).
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                                                      regulation; Activator; RNA-binding; Nuclear
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   10036 MW;
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72.7%;
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Pred.
   3832412849D5B1CE CRC64;
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No. 0.074;
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RESULT 11
TATLHVIA2
ID TATLH 10-00
AC P0461
DT 13-AU
DT 13-AU
DT 13-AU
DT 13-AU
DT 13-AU
DT TATLH 13-AU
DT 13-AU
TATL
RA Stemp
RA
RESULT 12

TAT_HV1C4

ID TAT_H
AC P0590

DT 01-NC
DT 16-OC
DT 16-OC
DT TAT P
GN TAT.
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Best Local Similarity
"---hes 8; Conserve
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Best Local S
Matches 8
                                             01-NOV-1988
01-NOV-1988
16-OCT-2001
                                                                                                                       TAT_HV1C4
P05907;
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P04614;
13-AUG-1987
13-AUG-1987
16-OCT-2001
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SEQUENCE
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MEDLINE-85090453; PubMed-2578227;

Sanchez-Rescador R., Power M.D., Barr

Stempien M.M., Brown-Shimer S.L., Gee

Levy J.A., Dina D., Luciw P.A.;
               TAT protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Science 227:484-492(1985).

11- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR001831; HIV_Tat. Pfam; PF00539; Tat; 1. PRINTS; PR00055; HIVTATDOMAIN
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
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SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
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(Rel. 09, Last sequence update)
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Pred. No. 0.08
3; Mismatches
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0.087;
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Renard A., Randolph A.,
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RESULT 13
TAT_HYLINE
AC P2087
AC P2087
DT 01-FE
DT 16-OC
DF TAT.
GN TAT.
GN TAT.
GN TAT.
CN VLrus
OX NCBL.
RN [1]
RP SEQUIT
CC -!-!
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Best Local
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01-FEB-1991
01-FEB-1991
16-OCT-2001
                                                                                                                                                                                         SEQUENCE FROM N.A.

KOYANA91 S., Chen I.S.Y.;

Submitted (DEC-1988) to the HIV data bank.

-I- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE L'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Molecular cloning and primary nucleotide sequence analysis of a distinct human immunodeficiency virus isolate reveal significant divergence in its genomic sequences.";

Proc. Natl. Acad. Sci. U.S.A. 83:8380-8384(1986).

- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO TRANS.ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THI
  the European
                                                    This SWISS-PROT entry is copyright. It is produced through
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=87041461; PubMed=3490666;
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                                                                                            PROMOTER.
SUBUNIT: BINDS CYCLIN T1 (BY SIMILARITY).
SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
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M13137; TAT$CDC45
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(Rel. 17, Last sequence update)
(Rel. 40, Last annotation update)
(Transactivating regulatory protein).
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Bioinformatics
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Best Local S
Matches 8
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Gurgo C., Guo H.-G., Franchini G., Aldovini A., Collalti E.,
Farrell K., Wong-Staal F., Gallo R.C., Reitz M.S. Jr.;
"Envelope sequences of two new United States HIV-1 isolates.";
"Involopy 164:531-536(1988).
"Involopy 164:531-536(1988).
"TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                POS905;
                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the El the European Bioinformatics Institute. There are no restruce by non-profit institutions as long as its content use by non-profit institutions as long as its content modified and this statement is not removed. Usage by anomaticity of the statement of the statement is not removed.
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                                                                                                                    entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL; M38429; -; NOT_ANNOTATED_CDS
HIV; M38429; TAT$JRCSF.
InterPro; IPR001831; HIV_Tat.
                                                   InterPro; IPR001831; HIV_Tat. Pfam; PF00539; Tat; 1.
                                                                              EMBL; M17449; AAA44851.1; -. HIV; M17449; TAT$MN.
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Viruses; Retroid viruses; Retroviridae; Lentivirus.
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16-OCT-2001
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SEQUENCE
                       Transcription
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ISOLATE WAS TAKEN I
                          Activator; RNA-binding; Nuclear
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Pred. No. 0.08
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Query Match Best Local S Matches 8

Similarity 8; Conser

Conservative

84.5%;

Score 49; DB Pred. No. 0.08
3; Mismatches

0.087;

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Length 101;

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RESULT 15

TAT_HV1OY

ID TAT_HV1OY

AC P20893;
DT 01-FEB-1991 (Rel. 17, Created)
DT 01-FEB-1991 (Rel. 17, Last sequence update)
DT 01-FCB-1991 (Rel. 40, Last annotation update)

Tat_wr protein (Transactivating regulatory protein).
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Search completed: July 15, 2002, 13:23:34 Job time: 1450 sec
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Matches 8; Conserv
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MEDLINE-90148544; PubMed-2559749;
Huet T., Dazza M.C., Brun-Vezinet F., Roelants G.E., Wain-Hobson S.;

"A highly defective HIV-1 strain isolated from a healthy Gabonese individual presenting an atypical western blot.";

AIDS 3:707-715(1989).

-i- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATISG TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LITR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro: IPR001831; HIV_Tat. Pfam; PR00539; Tat; 1. PRINTS; PR00055; HIVTATDOMAIN.
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-!- SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
-!- MISCELLANEOUS: THE OYI ISOLATE WAS TAKEN FROM THE BLOOD OF A
HEALTHY GABONESE INDIVIDUAL.
-!- MISCELLANEOUS: THE SINGLE C -> S SUBSTITUTION AT RESIDUE 22 OF THE
OYI TAT PROTEIN RENDERS IT INACTIVE, BUT MAY NOT ACCOUNT FOR THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
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47 YGRKKRRQRRR 57
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Pred. No. 0.087;
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MEDLINE-21322025; Pubmed-11429117;
HUSAIN M., Hahn T., Yedavalli V.R.K.,
"Characterization of HIV Type 1 tat S
Perinatal Transmission.";
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PROMOTER (BY SIMILARITY).
EMBL; ANO7059; AAG32106.1; -.
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PRINTS; PR00055; HIVTATDOMAIN.
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MEDLINE-21322025; PubMed-11429117;
Husain M., Hahn T., Yedavalli V.R.K.,
"Characterization of HIV Type 1 tat 5
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NCBI_TaxID=11676;
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-I- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
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FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATION RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LI
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AIDS Res. Hum. Retroviruses 17:765-773(2001).
-I- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
STRAIN-PAIRD, MOTHER;
MEDLINE-21322025; PubMed-11429117;
Husain M., Hahn T., Yedavalli V.R.K.,
                                                                                                      Viruses; Retroid viruses; NCBI_TaxID=11676;
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G REGULATORY PROTEIN) (FRAGMENT).
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2; Mismatches
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2; Mismatches
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AIDS Res. Hum. Retroviruses 17:765-773(2001).

1. FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR PROMOTER (BY SIMILARITY).

EMBL. AV007063; AAG32110.1;

InterPro; IPR00183; HY_Tat.

Pfam; PF00539; Tat; 1.

PFINTS; PR00055; HIVTATDOMAIN.
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NCBI_TaxID=11676;
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EMBL; AY007062; AAG32109.1; -
InterPro; IPR001831; HIV_Tat.
Pfam; PF00539; Tat; 1
PRINTS; PR00055; HIVTATDOMAIN.
                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                              Activator; Nuclear protein; RNA-binding; Transcription regulation NON_TER 72 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Husain M., Hahn T., Yedavalli V.R.K., Ahmad N. "Characterization of HIV Type 1 tat Sequences
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les 9; Conserv
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FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
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                                                                          PRELIMINARY;
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Best Local
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-!- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR PROMOTER (BY SIMILARITY).
EMBL: AV007067; AAG33214.1; -.
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STRAIN-PAIRD, MOTHER;
MEDLINE-21322025; PubMed=11429117;
HUSAIN M., Hahn T., Yedavalli V.R.K., Ahmad N.
"Characterization of HIV Type 1 tat Sequences
"Perinatal Transmission.";
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EMBL; AY007066; AAG32113.1; -
InterPro; IPR001831; HIV_Tat.
Pfam; PF00539; Tat; 1
PRINTS; PR00055; HIVTATDOMAIN.
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PRINTS; PR00055; HIVTATDOMAIN
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-i- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
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Viruses; Retroid viruses; Retroviridae; Lentivirus
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MEDLINE=21322025; PubMed~11429117;
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Sabino E., Pan L.Z., Cheng-Mayer C., Mayer A.;
Sabino E., Pan L.Z., Cheng-Mayer C., Mayer A.;
"Comparison of in vivo plasma and peripheral blood mononuclear cell
HIV-1 quasi-species to short-term tissue culture isolates: an analysis
of tat and C2-V3 env regions.";
AIDS 8:901-909(1994).
-i- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE
TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND
ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
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MEDLINE-21322025; PubMed-11429117;
HUSAIN M., Hahn T., Yedavalli V.R.K., Ahmad N.;
"Characterization of HIV Type 1 tat Sequences Associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Viruses; Retroid
NCBI_TaxID=11676;
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-I- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND
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"The TAT and C2-V3 Envelope Genes in the Molecular Epidemiolo
Human Immunodeficlency Virus-1.";
Submitted (MAY-1996) to the EMBL/GenBank/DDBJ databases.
1- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING
TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AN
ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM
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PRINTS; PR00055; HIVTATDOMAIN.
SEQUENCE FROM STRAIN-T10;
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STRAIN=PARTENT LTS1;
MEDLINE-99049627; PubMed-9833881;
Quinones-Mateu M.E., Mas A., Lain de Lera T., Soriano V., Alco
Lederman M.M., Domingo E.;
"LTR and tat variability of HIV-1 isolates from patients with
  Q9YP33
                                                                                                                                                                                                                              Lorenzo E., Herrera R.J., Lai S., Fischl M.A., Hill M.D.;
"The TAT and C2-V3 Envelope Genes in the Molecular Epidemiology of
Human Immunodeficiency Virus-1."
Submitted (MAY-1996) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE
TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND
ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
PROMOTER (BY SIMILARITY).
EMBL; U57282; AAB17842.1; -
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01-DEC-2001
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HIV-1 into South America.";
EMBL; AV037279; AAK12727.1; -. SEQUENCE 101 AA; 11525 MW; 52A6782BF08FAAB9 CRC64;
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Viruses; Retroid viruses; Retroviridae; Lentivirus
NCBI_TaxID=11676;
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Fang G., Weiser B., C
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Viruses; Retroid viruses; Retroviridae; Lentivirus
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Ol-MAY-1999 (TrEMBLrel. 10, Last sequence update)
Ol-DEC-2001 (TrEMBLrel. 19, Last annotation update
TAT PROTEIN (TRANSACTIVATING REGULATORY PROTEIN).
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1 YGRKKRRQKKK 11
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Interleukin 17 (hi Sequence deduced fi HIV-1 tat protein. TAT transduction d Tat peptide. Unid TAT protein transd Minimal eleven ami Beta-catenin deriv Peptide associated HIV TAT transducti
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					tor like protein; etic; immunosuppre; anti-alzhelmer's tic; dermatologica eukaemic; anti-inf a; neuronal dysfun der; eye disorder;	receptor like protein associated			11 AA.	ALIGNMENTS	AAV71012 AAV71012 AAU09932 AAU09812 AAE13064 AAE13064 AAE13064 AAE122605 AAE12204 AAB60221 AAB60226 AAB6025 AAB60266 AAB6045 AAB6045 AAB6045 AAB603418 AAB603418 AAB60373 AAE60373 AAE60373 AAE60373 AAE60373 AAE60373 AAE60373 AAE60373 AAE60373 AAE60376 AAE6037
					<pre>immunomodulatory; ssive; anti-microbial; , anti-parkinsonlan; l; renal; osteopathic; ertility; ophthalmological; ction; lung disease; cancer; human.</pre>	ated peptide #1.					Human immunodefici HIV-1 tat protein Protein transducti HIV-1 tat protein Protein transducti HIV-1 tat protein tr HIV-1 TAT protein tr HIV-1 TAT protein tr HIV-1 TAT peptide fici HIV tat protein tr Protein transducti HIV-1 TAT protein tr Protein transducti HIV-1 TAT protein tr HIV TAT protein tr Anti-herpetto pepti HIV-TAT protein tr Anti-herpetto pepti HIV-TAT protein tr Anti-herpetto fth inv HIV-TAT protein tr Anti-herpetto fth inv

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expression and activity.

Note: This artificial peptide sequence is given in the sequence listing how is not described in the specification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention describes novel nucleic acids encoding interleukin (IL) 17 receptor like polypeptides useful as vaccines and in gene therapy. These have immunomodulatory, anti-inflammatory, anti-diabetic, anti-microbial,
02-MAY-1990;
                                                                            15-NOV-1990
                                                                                                                                                                                                                                                            Region
                                                                                                                                                                                                                                                                                                                                 Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                     HIV 1-NDK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence deduced from tat gene of HIV 1-NDK.
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ygrkkrrgrrk 11
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    90WO-FR00312
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Pred, No. 0.04
2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        δõ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 an AIDS patient. A genomic library was prepd. from DNA extracted from CEM cells infected with the virus. The library was screened with a psm1 probe corresp. to a fragment from HIV 1. The virus is more cytopathic than other strains and is not inhibited by OKT4A. It has been deposited as CNCM I-857. The sequence can be used to express proteins useful for diagnosing the presence of NDK and related viruses and in vaccines against immunodeficiency diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New HIV-NDK retrovirus and protein component - used in against immuno-deficiency disorders and in raising MAbs retro-virus detection in vivo.
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N-PSDB; AAQ06635.
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                                                                                                                                                                                                                                                                                                                     HIV-1 tat protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Fig 2; 37pp; French
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Example 8; Page 85; 118pp; English
                      BH3 interacting domain death agonist polypeptide - used for treating decreased apoptotic conditions resulting from inflammation etc.
                                                                                                                                        09-SEP-1996;
                                                                                                                                                                  09-SEP-1997;
                                                                                                                                                                                                                   W09809980-A1
                                                                                                                                                                                                                                            Human immunodeficiency virus type
                                                                                                                                                                                                                                                                   Mouse; BH3 interacting domain death agonist; BID; BCL-2 family; apoptosis; regulation; cell death; inflammation; cancer; arthriautoimmune disease; viral infection; lymphoproliferative.
                                                                                                                                                                                                                                                                                                                                               20-JUL-1998 (first entry)
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                                                              WPI; 1998-193546/17.
                                                                                       Korsmeyer
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                                                                                                                                         96US-0706741
                                                                                                                                                                  97WO-US15872
                                                                                                                                                                                                                                                                                                                                                                                                  Protein;
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81.8%;
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Pred. No. 0.
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                                                                                                                                                                                                                                                                                cancer; arthritis;
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MAbs for
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        This invention describes a novel anti-pathogen system (APS) comprising a fusion protein constructed from a covalently linked protein transduction domain (PTD) and a cytotoxic domain. The APS can be used for suppressing
                                                                                                                                                          WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       treating a decreased apoptotic state of a cell. The decreased apoptotic state that is treated results from a disease such as cancer, viral infections, lymphoproliferative conditions, arthritis, inflammation and autoimmune diseases. Antibodies against the BID protein can be used for detecting a BID polypeptide in a cell or population of cell. The nucleic acid sequence and the BID protein can also be used for treating immunodeficiency disease (including AIDS), senescence, neurodegenerative disease, ischaemic and reperfusion cell death, infertility and wound-healing. Primers derived from the nucleic acid encoding the BID protein can be used for detecting/quantitating the protein and for detecting alterations in the nucleic acid encoding the BID protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Anti-pathogen; fusion protein; protein transduction domain; PTD; AZT; cytocoxic domain; suppressor; infection; medicament; ddl; ddC; d4T; 3TC; FTC; DAPD; 1592089; CS92; acyclovir; anciclovir; peniclovir; interferon; apoptosis; virus; HIV; cytomegalovirus; CMV; herpes simplex virus; HSV-1; hepatitis virus; Kaposi's sarcoma-associated herpes virus; KSHV; herpes virus; yellow fever virus; flavivirus; rhinovirus; plasmodial;
                                                                                                                                                                                          Dowdy
                                                                              Claim
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10-DEC-1997;
                                                                                                                                                                                                                                                                                                                                    17-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                  Unidentified
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                                                                                                                        New anti-pathogen systems, particularly for virus
                                                                                                                                                                                                                        VIND ( WIND)
                                                                                                                                                                                                                                                                                                     10-DEC-1998;
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                                                                           65; Page
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                                                                           34; 123pp; English
                                                                                                                                                                                                                                                       98US-0082402
97US-0069012
                                                                                                                                                                                                                                                                                                     98WO-US26358
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72.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cc administering a medicament e.g. AZT, ddI, ddC, d4T, 3TC, FTC, DAPD, CC 1592U89, CS92, acyclovir, ganciclovir, peniclovir or an interferon. The CC APS can also be administered to a mammal in the presence of a pathogen to CC induce apoptosis in a predetermined population of cells. The products can CC be used for treating mammals suffering from or susceptible to a viral CC infection or a disease associated with a virus, e.g. HTV, cytomegalovirus (CC (CMV), herpes simplex virus, e.g. type 1 (HSV-1) hepatitis virus, type CC (HCV), Kaposi's sarcoma-associated herpes virus (KSHV or human herpes CC (HCV), Kaposi's sarcoma-associated herpes virus (KSHV or human herpes CC (HCV), Kaposi's sarcoma-associated herpes virus (KSHV or human herpes CC (HCV), Kaposi's sarcoma-associated herpes virus (KSHV or human herpes CC (HCV), Kaposi's parcoma-associated herpes virus (KSHV or human herpes CC (HCV), Kaposi's parcoma-associated herpes virus (KSHV or human herpes CC (HCV), Kaposi's parcoma-associated herpes virus (KSHV or human herpes CC (HCV), Kaposi's parcoma-associated herpes virus, or suffering from CC (HCV), Kaposi's parcoma-associated herpes virus (KSHV or human herpes CC (HCV), Kaposi's parcoma-associated herpes virus, or suffering from CC (HCV), Kaposi's parcoma-associated herpes virus, or suffering from CC (HCV), Kaposi's parcoma-associated herpes virus, or suffering from CC (HCV), Kaposi's parcoma-associated herpes virus, or suffering from CC (HCV), Kaposi's parcoma-associated herpes virus, or suffering from CC (HCV), Kaposi's parcoma-associated herpes virus, or suffering from CC (HCV), Kaposi's parcoma-associated herpes virus, or suffering from CC (HCV), Kaposi's parcoma-associated herpes virus, or suffering from CC (HCV), Kaposi's parcoma-associated herpes virus, or suffering from CC (HCV), Kaposi's parcoma-associated herpes virus, or suffering from CC (HCV), Kaposi's parcoma-associated parcoma-associated herpes virus, or suffering from CC (HCV), Kaposi's parcoma-associated herpes virus, or suffering from CC (HCV)
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Best Local
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This sequence represents a Tat peptide. The invention relates to a bel homology domain 3 (BH3 domain), derived from a proapoptotic member of the BCL-2 family. The BH3 polypeptide can be used in a method for promoting apoptosis target cell, especially where the cell is a cancer cell a virus cell or an autoantibody producing cell. The BH3 polypeptide can
                                                                                                                                                                                                                                                                            WPI; 1999-255058/21
                                                                                                                                                                                                                                                                                                                                                                                                                       07-OCT-1997;
26-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BH3 domain; cell death agonist; bcl homology domain; BCL-2 family; apoptosis promoter; cancer cell; virus infected cell; inflammation; autoantibody producing cell; cancer; lymphoproliferative condition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY05415 standard; peptide; 11 AA
                                                                                                                                                                         Claim 7; Page 67; 104pp; English.
                                                                                                                                                                                                                        Bcl homology domain 3 polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W09916787-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            arthritis; autoimmune
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY05415;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pathogen strains. This sequence represents a TAT transduction domain peptide motif which is used in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                          WASHINGTON
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                                                                                                                                                                                                                                                                                                                                                                                                                       97US-0946039.
97US-0060133.
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72.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 49; DB 2
Pred. No. 0.12;
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be used
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RESULT AAABOLT AAABOLT
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Best Local
                                                   beta-catenin are described. The modulating agent comprises an internalisation modety and one or more of an amino acid sequence sylds(PO_4)GIHS(PO_4)G. or a peptide analogue or peptideminetic of the manufacture of a medicament for stimulating agents are useful for the manufacture of a medicament for stimulating hair growth or reducing hair loss, stimulating skin exfoliation, and inhibiting the development of Alzhaimer's disease. They may also be used to increase the beta-catenin level in a cell, to stimulate activation of gene transcription in a cell, and to stimulate cell differentiation. They may further be used to ameliorate hearing loss resulting from a variety of inner ear disorders, such as hyperacusis and tinnitus, through regeneration of hair cells of the inner ear.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         in therapeutic compositions for treating disease including cancer, other lymphoproliferative conditions, arthritis, inflammation, and autoimmune diseases, which may result from the down regulation of cell death
Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                           Modulating agents for inhibiting degradation of cytoplasmic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; Page 46; 49pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Modulating agents for inhibiting degradation of cytoplasmic beta-catenin, used for e.g. stimulating hair growth or reducing loss, inhibiting development of Alzheimer's disease, comprise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ₩PI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ADHE-) ADHEREX TECHNOLOGIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus (HIV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Modulating agent; beta-catenin; hair loss; hair growth; skin;
exfoliation; Alzhelmer's disease; gene transcription;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TAT protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                internalization moiety and
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les 8; Conser
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      11
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72.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                amino acid sequence
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Pred. No. 0.12
3; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inner ear; hyperacusis; tinnitus;
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Query Match

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                                        shutting off the downstream Fas signalling pathway. The chimeric group and fusion peptide are useful for inhibiting ligand-induced apoptosis by bringing them into contact with T cells. The chimeric group is useful for expanding T cells in vitro e.g. T cells specific for particular antigens such as tumour-specific antigen, for enhancing immune response and to inhibit the apoptosis of chronically activated T cells e.g. activated CD4^+ T cells in HIV infected patients. The chimeric group is also useful for therapeutic, prophylactic or diagnosis of intracellular delivery of small molecules and macromolecules such as anti-spoptotic polypeptides and nucleic acids encoding such polypeptides. Two primers (AAA54297, AAA54298) were used to amplify the FLIP CDNA for subcloning into the XhoI-NcoI site of the pHA-TAT vector which contains the N-terminal protein translator of the properties from the human impunded ficiency virus tat
                                                                                                                                                                                                                                                                                                                                                                 across the cell membrane. The anti-apoptotic polypeptide is FUICE-like inhibitor protein (FLIP) which inhibits Fas and TNF mediated apoptosis by inhibiting binding of Caspase-8 to the Fas receptor complex, thus
                                                                                                                                                                                                                                                                                                                                                                                                                                            A chimeric group or fusion peptide which comprises a portion of an anti-apoptotic polypeptide which inhibits apoptosis of lymphocytes combination with a transport group is described. The transport group is capable of transporting the chimeric group or fusion peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           group
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06-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chimeric protein;
FLIP; Fas; TNF; a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fusion polypeptide comprises portion of
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  protein.
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8; Conserv
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Tat is a preferred
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 30;
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99US-0128021
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ㅁ
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of anti-a
                             from the human immunodeficiency virus tat
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Best Local
                                                                                                                                             The present invention is concerned with methods of modulating the amount of free beta-catenin in the cell, and methods of stimulating the expression of genes involved in cellular differentiation, the transcription of which is under the control of beta-catenin. The peptides given in AAB27053-B27088, AAB27284-P27300 and AAB27330-B27351 can be used as modulating agents which interrupt the interaction between alpha and beta catenin, causing increased levels of the latter and stimulating the activation of beta-catenin mediated transcription. This can be used to stimulate cell differentiation, which can then be used to promote hair growth and skin exfoliation. This latter is particularly useful in the improvement of photodamaged skin and to minimise wrinkles. The modulating peptide can also be used to reduce hearing loss resulting from inner ear disorders such as hyperacusis and tinnitus.
                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                 Stimulating beta-catenin mediated gene expression, cellular differentiation and hair growth, involves contacting cells with modulating agent capable of inhibiting interaction between alpha beta catenin
                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 19; 77pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Beta-catenin; cadherin-mediated intercellular adhesion; cell differentiation; modulating agent; hair loss; skin exfoliation; internalisation molety; flanking sequence; transcription; hearing lo
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ygrkkrrqrrr 11
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72.7%;
                                                    Score 49; DB Pred. No. 0.1; 3; Mismatches
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Pred. No.
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Protein transduction domain; fusion molecule; therapeutic agent; drug targetting; drug discovery; cell transduction; bioavailability; vaccine; nervous system disorder; Alzheimer's disease; Parkinson's disease; Huntington's disease; pre-senile dementia; epiloseizure; compulsive behaviour; meningitis; encephalitis; ischaemia;
                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a method for modulating beta-catenin mediated gene transcription in a cell. The method involves contacting a cell with a modulating agent comprising an internalization modety and a peptide comprising a sequence LXXLL or peptide analogue of LXXLL. The method is useful for modulating beta-catenin mediated gene transcription, cell differentiation, hair growth, and retinoic acid activity, for treating cancer, and for inhibiting the development of
                                                                              HIV TAT
                                                                                                     09-FEB-2001
                                                                                                                             AAB29413;
                                                                                                                                                    AAB29413 standard;
                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                      Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Use of modulating agent comprising internalization moiety and peptide, for modulating beta-catenin mediated gene transcripts cell differentiation, for treating cancer, and for inhibiting Alzheimer's disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-679589/66.
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cancer; Alzheimer's
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                                                                             transduction
                                                                                                                                                                                                                                                                  Similarity
8; Conser
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                                                                                                                                                    peptide;
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                                                                                                                                                                                                                                                                             84.5%;
                                                                           domain,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English.
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                                                                                                                                                                                                                                                                            Score 49; DB 2
Pred. No. 0.12;
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epilepsy;

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RESULT , 11
AAB09907
ID AAB099
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                                                                                                                                                                                                        The invention relates to a novel fusion molecule comprising at least CC one protein transduction domain (PTP) and at least one linked molecule, where the linked molecule has therapeutic or prophylactic activity CC against a medical condition. The invention also relates to methods of CC drug discovery in which the test compound is linked to a sultable CC transducing protein and introduced to a cell; a method of killing CC comprising a covalently linked fusion molecule; a mammal adapted for CC experimental use in which at least one transduction molecule; a mammal compound is linked fusion molecule; a mammal adapted for CC transduced into essentially all the cells of the mammal. The fusion CC molecule is used to deliver a therapeutic agent to a mammal, especially CC a human. The linked molecule may be a vaccine, an anti-infective drug, CC a cardiovascular drug, an antitumour drug, an analyssic, an CC entral nervous system (CNS) disorder is especially Alzheimer's disease. CC central nervous system (CNS) disorder is especially Alzheimer's disease, (including viral and bacterial meningitis), encephalitis, ischaemia, CC (including viral and bacterial meningitis), encephalitis, ischaemia, CC (used to kill virally infected cells, especially those infected with HIV. CC The wackings are used to treat or prevent bacterial or viral infections. The methods are a highly effective means for transducing a molecule collent transduction of the target cell. The problems that transduction of the target cell. The present sequence cC hydrophobicity, hydrophilicity, biological resistance), by providing CC efficient transduction of the target cell. The present sequence
                                                                                                                           Matches
                                                                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         spongiform encephalopathy; dyslexia; age-related memory loss; Lou Gehring's disease; viral infection, HIV; bacterial infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 50; 191pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fusion molecules comprising protein transduction domains therapeutic agents, useful for treating e.g. Alzheimer's Parkinson's diseases, dementia and epilepsy .
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29-AUG-1999;
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          AAB09907 standard; peptide;
                                                                                                                                                                                       Sequence
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99US-0151291
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72.7%;
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Pred. No. 0
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RESULT 12
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Best Local S
Matches 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disease, Reiter's disease, Celiac disease, sympathetic ophiculatura and primary biliary cirrhosis, immune disorders, graft versus host disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Use of agent capable of intracellularly inhibiting mucin MUC-1 for inducing T-cell-based immunosuppression and for treating autoimmune disorders, transplant rejection and inflammatory disorders .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI;
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immune disorder; inflammatory disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV tat protein transduction
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Agrawal B,
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                                                                                                                                                                                                                                                                                                                                    AAY93542;
                                                                                                                                                                                                                                                                                                                                                                                                                      AAY93542 standard; Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JUN-2000
cytotoxic domain; pathogen infection;
plasmodial infection; cancer; prostate
                                                                                  Protein transduction system; protein transduction domain;
                                                                                                                                                                Amino acid sequence of a synthetic protein transduction domain
                                                                                                                                                                                                                                                  25-SEP-2000
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| ygrkkrrqrrr ||
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8; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11
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72.78;
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Pred. No. 0.12;
prostate
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                                             retroviral infection;
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RESULT 13
AAYT10
XX AAYT10
ACC AAYT10
XX 29-AUG
XX 29-AUG
XX Phosph
KW Phosph
KW TAT pr
KW Cardia
XX Human
XX H
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Best Local
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                                                                                                                                                                                                                                                                                                                                            Phospholamban; PLB; sarcoplasmic reticulum Ca 2+ ATPase; SERCA2a; HIV; TAT protein; penetratin; transport peptide; cardiomyocyte; inhibitor; cargo peptide; contractilin; cardiac contractility; cardiant; treatment;
                          02-NOV-1998;
27-JUL-1999;
                                                                                                          02-NOV-1999;
                                                                                                                                                                  11-MAY-2000
                                                                                                                                                                                                                    WO200025804-A2
                                                                                                                                                                                                                                                                      Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human immunodeficiency virus TAT peptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY71015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY71015 standard; peptide; 11 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protein transduction system for treating cancer and pathogenic infections has a fusion protein comprising a protein transduct: domain covalently linked to a cytotoxic domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 10;
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                                                                                                                                                                                                                                                                                                                           disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          h 84.5%;
Similarity 72.7%;
8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11
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                          98US-0106718
99US-0145883
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                                                                                                                                                                                                                                                                                                                              heart
                                                                                                                                                                                                                                                                                                                        failure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 49;
Pred. No.
                                                                                                                                                                                                                                                                                                                        myocardial dysfunction
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0.12;
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RESULT 14
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Best Local
                           WPI; 2001-596910/67
                                                      Jing S,
                                                                                                               13-MAR-2000; 2000US-188786P
                                                                                                                                                                        20-SEP-2001
                                                                                                                                                                                                                                 Human
                                                                                                                                                                                                                                                         Human; fibroblast growth factor-like; FGF-L; HIV tat protein; protein transduction domain; PDT; human immunodeficiency viru
                                                                                                                                                                                                                                                                                                                                                                                             AAU09932 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Treating cardiac diseases, e.g. heart failure or myocardial dysfunction comprises enhancing cardiac contractility by inhibiting interaction between phospholamban and sarcoplasmic reticulum calcium 2+ adenosine
                                                                                    (AMGE-)
                                                                                                                                          13-MAR-2001; 2001WO-US08013
                                                                                                                                                                                                     WO200168854-A2
                                                                                                                                                                                                                                                                                                          Human
                                                                                                                                                                                                                                                                                                                                     15-JAN-2002
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                                                                                                                                                                                                                                                                                                    immunodeficiency virus (HIV) tat partial peptide sequence
                                                                                                                                                                                                                            immunodeficiency virus type
                                                                                    AMGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity 72. 8; Conservative
                                                       Bass MB;
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Wang Y, Silverman GJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11 AA;
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                                                                                  INC
                                                                                                                                                                                                                                                                                                                                                                                             Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  56pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            84.5%;
72.7%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 21;
                                                                                                                                                                                                                                                           immunodeficiency virus
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New fibroblast growth factor-like polypeptide and polynucleotide for

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RESULT 15
AAU09912
ID AAU099
XX 27-FEE
XX 27-FEE
XX 27-FEE
XX 61brob
KW 1mmunc
KW 005teoo
KW hyperr
KW haemoo
KW hyperr
KW human
XX hyperr
XX 27-SEI
PA 22-MA
XX 22-MA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC factor-like (FGF-L) nucleit acid molecules and polypeptides. The FGF-L cc polypeptide is useful for determining whether a compound inhibits FGF-L coplypeptide. The FGF-L polypeptides of the invention are useful for ct treating, preventing or ameliorating a compound that binds to the cc polypeptide. The FGF-L polypeptides of the invention are useful for cc treating, preventing or ameliorating a medical condition or an FGF-L coplypeptide-related disease, condition or disorder such as wound healing cd disorders, ulcers, gut disorders, lung disorders, liver disorders such as compatitive and disbetes. The invention is also useful for diagnosing a cc hepatitis and disbetes. The invention is also useful for diagnosing a cc ambject and is useful for modulating levels of FGF-L polypeptide and for identifying or quantifying the amount of FGF-L polypeptide and for identifying or developing novel agonists and antagonists of the cc or more diseases or disorders, and also as an immunogen for producing cantibodies for in vivo imaging. The present sequence represents an il contact the contact the contact of the contact of the contact the contact of t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          fibroblast growth factor receptor-like protein; FGFR-L; anorectic; haemostartic; osteopathic; cytostatic; nephrotropic; antidiabetic; immunomodulator; antiinflammatory; haematopoietic disorder; osteoporosis; osteogenesis imperfecta; Paget's disease; periodontal disease; cancer; hypercalcaemia; acute glomerulonephritis; chronic glomerulonephritis; diabetes; obesity; cachexia; transgenic animal; gene therapy;
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Novel nucleic acid encoding fibroblast growth factor receptor-like polypeptides, useful for treating hematopoietic disorder, osteopore paget's disease, glomerulonephritis, cancer, diabetes, obesity and
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                                                                                                                                                                                   osteoporosis,
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Disclosure; Page 59; 163pp; English

CC hypercal caemia, acute glomerulonephritis, chronic glomerulonephritis, contact context, and abbetes, obesity and cachexia. (1) is also useful for centerifying a compound which binds to FGFR-L polypeptide, by contacting CC (1) with a compound, determining the extent of binding of the FGFR-L CC polypeptide to the compound, and determining the activity of the polypeptide in an animal. A transgenic animal comprising (1) CC is useful for determining whether a compound inhibits FGFR-L polypeptide cativity or FGFR-L polypeptide production, by exposing the transgenic cativity or FGFR-L polypeptide production, by exposing the transgenic continuity of FGFR-L polypeptide production, by caposing the transgenic continuity of FGFR-L polypeptide production in CC the mammal. (II) is useful for mapping the locations of FGFR-L gene and CC essays to test for the presence of an FGFR-L nucleic acid molecule in CC mammalian tissue or bodily fluid samples, in gene therapy, and as tools CC immunogen, and for cloning FGFR-L polypeptide genes. (I) is useful as CC expression cloning strategy. The present sequence represents the amino acid sequence of human immunodeficiency virus (HTV-1) tat protein CC transduction domain (TAT PDT) as described in the invention. The invention relates to a novel isolated fibroblast growth factor receptor-like (FGFR-L) polypeptide (I). (I) and the nucleic acid (II) encoding (I) are useful for treating, preventing or ameliorating a medical condition including haematopoletic disorder, osteoporosis, osteogenesis imperfecta, Paget's disease, periodontal disease,

Sequence 11 Ā

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Matches
       Query Match
Best Local
Local Similarity
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Score 49; DB
Pred. No. 0.12
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Search completed: July 15, Job time: 415 sec 2002, 12:57:59

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re greater than or equal to the score of the result being printed,
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US-08-235-403-47
US-08-235-6077-4
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PCT-US95-06077-12
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2 2, Appl
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50, Appl
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65, Appl
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1 Y	ery Mat st Loca tches	RESULT 1 WS-08-706-741B-54 WS-08-706-741B-54 Sequence 54, Application US/08706; Patent No. 595593 Patent No. 595593 Patent No. 595593 Patent No. 595593 GENERAL INFORMATION: BH3 INTER MUNBER OF INVENTION: BH3 INTER CORRESSE: HOWELL & HAFERKAN STREET: 7733 FORSYTH BLVD., CITY: ST. LOUIS STATE: MISSOURI COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC COMPATIBLE POSYMS- SOFTWARE: Patentin Release # CURRENT APPLICATION NUMBER: US/08/76 FILING DATE: 09-SEP-1996 CLASSIFICATION INFORMATION: NAME: HOLLAND, DONALD R REGISTRATION NUMBER: 35,197 REFERENCE/DOCKET NUMBER: 965 TELEPHONE: (314) 727-5188 TELEPHONE: (314) 727-5188 TELEPRATION FOR SED ID NO: 54: SEQUENCE CHARACTERISTICS: LENGTH: 11 maino acids TOPOLOGY: Linear MOLECULE TYPE: peptide US-08-706-741B-54		28 32 32 32 33 33 34 34 44 44 44 44 44 44 44 44 44
YGRKKRRQKKK ::: YGRKKRRQRRR	h Simi	1706-741B-54 rence 54, Application rence 595593 legal INFORMATION: FPLICANT: KORSMEYER TILE OF INVENTION: UNBER OF SEQUENCES: OUNDERS PONDERCE ADDRESSE: HOWELL STATE: 7733 FORSY CITY: ST. LOUIS COUNTRY: USA ZID: 63146 MEDIUM TYPE: Flopp COMPUTER: READABLE FOR MEDIUM TYPE: Flopp COMPUTER: IBM PC OPERATING SYSTEM: SOFTWARE: PACENTION UNMERR: FILING DATE: 09-SE CLASSIFICATION UNMERR: FILING DATE: 09-SE CLASSIFICATION UNMERR: FILING DATE: 09-SE CLASSIFICATION UNMERR: FILING DON REGISTRATION UNMERR: FILEPHONE: (314) 7 TELEPHONE: 11 amino acid STRANDEDNESS: TOPOLOGY: Linear OLECULE TYPE: pepti706-741B-54		4444444444444444 000000000000000000000
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211	lty servat	AMATION: BH3 IN SOQUENCES: 88 3 IN SOQUENCES: 88 5 FLORDY disk EFLORDY disk EFLORDY disk IBM PC compatit SYSTEM: PC-DOS PATENTIN PC-DOS PATENTIN PC-DOS PATENTIN DATA: 109 - SEP-1996 (VION: 5.14 NUMBER: 35. TON NUMBER: 35		ហាហាហាហាហាហាហាហាហាហាហាហាហាហាហាហាហាហាហា
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US-08-924-695A-54

Sequence 54, Application US/08924695A Patent No. 5998583

GENERAL INFORMATION:

APPLICANT:

KORSMEYER,

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                                                                                                                                    Sequence 2, Application US/09208966

Patent No. 6221355

GENERAL INFORMATION:
APPLICANT: DOWDY, Steven F.
TITLE OF INVENTION: ANTI-PATHOGEN SYSTEM AND METHODS OF USE THEREOF FILE REFERENCE: 48881/1742

CURRENT APPLICATION NUMBER: US/09/208,966

CURRENT FILING DATE: 1998-12-10

EARLIER APPLICATION NUMBER: 60/082,402

EARLIER APPLICATION NUMBER: 60/089,012

EARLIER FILING DATE: 1998-04-20

EARLIER FILING DATE: 1998-04-20

EARLIER FILING DATE: 1998-04-20

EARLIER FILING DATE: 1997-12-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 35,197
REFERENCE/DOCKET NUMBER: 971798
TELECOMMUNICATION INFORMATION:
TELEPHONE: (314) 727-5188
TELEPHONE: (314) 727-6092
INFORMATION FOR SEO ID NO: 54:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPER: Amino acids
                                                        NUMBER OF SEQ ID NOS: 57
SOFTWARE: PatentIn Ver. :
SEQ ID NO 2
LENGTH: 11
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Best Local Similarity
             TYPE: PRT
ORGANISM: human
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PALENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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ATTORNEY/AGENT INFORMATION:
NAME: HOLLAND, DONALD R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
TOPOLOGY: linear
MOLECULE TYPE: peptide
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STATE: MIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/924,695A FILING DATE: 09-SEP-1997
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7733 FORSYTH BLVD., SUITE 1400
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H3 INTERACTING DOMAIN DEATH AGONIST
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Pred. No. 0.067;
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; ORGANISM: Human immunodeficiency virus
US-09-296-089-37
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SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 37
LENGTH: 11
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Matches 8; Conserv
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CURRENT APPLICATION NUMBER: US/09/296,089
CURRENT FILING DATE: 1999-04-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Blaschuk, Orest W.
APPLICANT: Byers, Stephen
APPLICANT: Gour, Barbara J.
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: FISH & NE
                                                               FILING DATE: 25-MAY-
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                   STREET: LL. STREET: New York CITY: New York TISA
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TITLE OF INVENTION: TAT-DERIVED TRANSPORT POLYPEPTIDES
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nes 8; Conserv
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APPLICATION NUMBER: US 0
FILING DATE: 28-APR-1994
APPLICATION NUMBER: US 0
FILING DATE: 21-AUG-1992
                                                                                                                                                                 OPERATING SYSTEM: PC-DOS/MS-DOS
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o. 5652122
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                US 07/934,375
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                                                US 08/235,403
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Pred. No. 0.067;
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Pred. No. 0.067;
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19-AUG-1993

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US-08-450-246-47
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Patent No. 5
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Best Local :
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TELEX: 14-8367
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM CCO-compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/235,403
FILING DATE: 28-APR-1994
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TITLE OF INVENTION: TAN DERIVED TRANSPORT POLYPEPTIDES
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                                                                                                                        APPLICATION NUMBER: FILING DATE: 25-MAN CLASSIFICATION: 51
            APPLICATION NUMBER: US 07/934,375
FILING DATE: 21-AUG-1992
APPLICATION NUMBER: US 07/098,766
FILING DATE: 28-JUL-1993
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nes 8; Conserv
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TYPE: amino acid
 APPLICATION NUMBER:
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STREET: 1251 Avenue of the Americas
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FILING DATE: 21-DEC-1989
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REFERENCE/DOCKET NUMBER: B170 CIP
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FILING DATE: 02-JAN-1991
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PCT/US93/07833
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Pred. No. 0.072;
3; Mismatches
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RESULT 7
US-08-450-098-47
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APPLICANT: FRANKEL
APPLICANT: PABO, C
APPLICANT: BARSOUM
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                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: BARSOUM, James G.

APPLICANT: FRMELL, Stephen E.

APPLICANT: PEPINSKY, R. B.

TITLE OF INVENTION: TAT-DERIVED TRANSPORT POLYPEPTIDES

NUMBER OF SEQUENCES: 69
                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/235,403
FILING DATE: 28-APR-1994
                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
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FILING DATE. US U///--,
APPLICATION NUMBER: US U///--,
FILING DATE: 28-JUL-1993
APPLICATION NUMBER: PCT/US93/07833
APPLICATION NUMBER: 19-AUG-1993
"C 07/454,450
                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                     APPLICATION NUMBER:
FILING DATE: 25-MAN
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nes 8; Conserva
                                                                                   APPLICATION NUMBER: US 07/934,375
FILING DATE: 21-AUG-1992
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STRANDEDNESS: sin
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25-MAY-1995
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Pred. No. 0.072;
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APPLICATION NUMBER: US 08/158,015 FILING DATE: 24-NOV-1993

FILING DATE: APPLICATION NUMBER:

пыльки: US 07/636,662 02-JAN-1991

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; TOPOLOGY: linear; MOLECULE TYPE: peptide US-08-450-098-47
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Best Local Similarity
% 8; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 47
                                                                               APPLICATION NUMBER: US/08/451,233
FILING DATE: 25-MAY-1995
CLASSIFICATION 514
PRIOR APPLICATION DATA: 08/235,403
APPLICATION NUMBER: US 08/235,403
FILING DATE: 28-APR-1994
APPLICATION NUMBER: US 07/934,375
FILING DATE: 21-AUG-1992
APPLICATION NUMBER: US 07/938,766
FILING DATE: 28-JUL-1993
APPLICATION NUMBER: PCT/US93/07833
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (212) 596-901
TELEFAX: (212) 596-9090
TELEY 14-8367
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 24-NOV-1993
ATTORNEY/AGENT INFORMATION:
NAME: Haley Jr., James F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: B170 CIP 2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 12 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Haley Jr., James F. REGISTRATION NUMBER: 27,7
                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 1251 Avenue of the Americas CITY: New York
                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
 APPLICATION NUMBER:
                                   APPLICATION NUMBER:
                                                                   APPLICATION NUMBER:
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linear
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VENTION: TAT-DERIVED TRANSPORT POLYPEPTIDES
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                                                   21-DEC-1989
                   02-JAN-1991
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                                                                     US 07/454,450
                                 US 07/636,662
US 08/158,015
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Pred. No. 0.072;
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Matches 8; Conserv
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TELEFAX: (212) 596-9090
TELEX: 14-8367
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08, FILLING DATE: 25-MAY-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08, FILING DATE: 28-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: B170 CIP
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: FRANKEL, Alan
APPLICANT: PABO, Carl
APPLICANT: BARSOUM, James G.
APPLICANT: FAMELL, Stephen E.
APPLICANT: PEPINSKY, R. B.
TITLE OF INVENTION: TAT-DERIVED TRANSPORT POLYPEPTIDES
NUMBER OF SEQUENCES: 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 24-NOV-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: FISH & NEAVE
ATTORNEY/AGENT
NAME: Haley
                      APPLICATION NUMBER: US 08
APPLICATION NUMBER: US 08
APPLICATION 24-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY
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                                                                                                                                                                                                       APPLICATION NUMBER: US 07/934,375 FILING DATE: 21-AUG-1992
                                                                     APPLICATION NUMBER: US 07/636,662 FILING DATE: 02-JAN-1991
                                                                                                        FILING DATE:
                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                       APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
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linear
                  INFORMATION:
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25-MAY-1995
                                                                                                                                      19-AUG-1993
                                                                                                   21-DEC-1989
                                                                                                                                                                        28-JUL-1993
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                                                                                                                                                                                        US 07/098,766
                                                                                                                     US 07/454,450
                                                                                                                                                         PCT/US93/07833
                                                                                                                                                                                                                                                              US 08/235,403
                                                  US 08/158,015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             47:
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Pred. No. 0.072;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 12;
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; TOPOLOGY: linear; MOLECULE TYPE: peptide US-08-733-505A-50
              QУ
                             Query Match
Best Local Similarity
Watches 8; Conserve
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US-08-733-505A-50
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Best Local
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                                                                                                                                                                                                                            TELEFAX: (314) 727-6092
INFORMATION FOR SEQ ID NO: !
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: KORSME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: HOLLAND, DONALD R.
REGISTRATION NUMBER: 35,197
REFERENCE/DOCKET NUMBER: 96:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE:
CLASSIFICATION: 530
                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: KORSMEYER, STANLEY J.

TITLE OF INVENTION: SERINE SUBSTITUTED MUTANTS OF

TITLE OF INVENTION: BCL-XL/BCL-2 ASSOCIATED CELL DEATH REGULATOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: B170 CIP 2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 7733 FO
CITY: ST. LOUIS
STATE: MISSOURI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER:
                                                                                                                                                              STRANDEDNESS
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nes 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 63105
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1 YGRKKRRQKKK 11
                                                                                                                                                                                                                                                        TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: HOWELL & HAFERKAMP, L.C.
STREET: 7733 FORSYTH BLVD., SUITE 1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
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                                            Conservative
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                                                          84.5%;
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72.7%;
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                                                                                                                                                                                                                                                                                      965458
                                                         Score 49; DB 2;
Pred. No. 0.072;
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Pred. No. 0.072;
                                             Mismatches
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                                          0;
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                                                                      Length 12;
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                                          Indels
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                                          0;
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                                                Query Match
Best Local Similarity
Watches 8; Conserva
                                                                                                                                                   ; TOPOLOGY: linear; MOLECULE TYPE: peptide US-08-235-403-47
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US-08-235-403-47
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                                                                                                                                                                                                                                                 TELEFAX: (212) 596-9090
TELEX: 14-8367
INFORMATION FOR SEO ID NO: 4
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07,
FILING DATE: 21-AUG-1992
APPLICATION NUMBER: US 07,
FILING DATE: 28-UUL-1993
APPLICATION NUMBER: PCT/US
FILING DATE: 19-AUG-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 27,794
REGERENCE/DOCKET NUMBER: B170 CIP 2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: BARSOUM, James G.
APPLICANT: FAWELL, Stephen E.
APPLICANT: PEPINSKY, R. B.
TITLE OF INVENTION: TAT-DERIVED TRANSPORT POLYPEPTIDES
                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                          NAME: Haley Jr., James REGISTRATION NUMBER: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 1251 Ave
CITY: New York
STATE: New York
                                                                                                                                                                                                                          TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 07/636,662
FILING DATE: OS-JAN-1991
APPLICATION NUMBER: US 08/158,015
FILING DATE: 24-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 07/454,450 FILING DATE: 21-DEC-1989
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                             1 YGRKKRRQKKK 11
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YGRKKRRQRRR 11
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                                                                   Conservative
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                                                                                Score 49; DB 4;
Pred. No. 0.072;
                                                                 Mismatches
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RESULT 12 PCT-US95-06077-4

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Best Local (
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INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
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REFERENCE/DOCKET NUMBER: IR
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9200
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COMPUTER READABLE FORM:
MEDIM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERAGING SYSTEM: PC-DOS/MS-DOS
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TITLE OF INVENTION:
TITLE OF INVENTION:
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APPLICATION NUMBER: US 06
FILING DATE: 23-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
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MEDIUM TYPE: Floppy disk
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: 'PatentIn Release #1.0, Version #1.25
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CITY: Spring House
STATE: Pennsylvania
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res 8; Conserv
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                                                                                        COUNTRY:
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VENTION: Vaccine Interdiction of Extracellular
VENTION: Transactivating Proteins of Human Immunodeficiency Virus
VENTION: and Other Chronically Infecting Viruses Employing Similar
VENTION: Intercellular Transactivating Strategies
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23-MAY-1994
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Transactivating Proteins of Human Immunodeficiency Virus
and Other Chronically Infecting Viruses Employing Similar
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Pred. No.
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                   TELEFAX: 215-540-5818
INFORMATION FOR SEQ ID NO:
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ATTORNEY/AGENT INFORMATION:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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TITLE OF INVENTION:
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                 SEQUENCE CHARACTERISTICS
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER:
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TELEPHONE: 215-540-5818
TELEPHONE: 215-540-5818
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                                                                                                                         NAME: Bak, Mary E. REGISTRATION NUMBER:
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                                                                      TELEPHONE:
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Spring House
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Pred. No. 0.081;
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Search completed: July 15, 2002, 12:59:20 Job time: 391 sec
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PCT-US95-06077-12
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; MOLECULE TYPE: peptide
PCT-US95-06077-11
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PCT-US95-06077-12
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GENERAL INFORMATION:
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Best Local Similarity
Matches 8; Conserv
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TELEFAX: 215-540-9818
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: IR:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9200
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/06077
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/247,991
FILING DATE: 23-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
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MEDIUM TYPE: Floppy disk
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TITLE OF INVENTION: Vaccine Interdiction of Extracellular
TITLE OF INVENTION: Transactivating Proteins of Human Immunodeficiency Virus
TITLE OF INVENTION: and Other Chronically Infecting Viruses Employing Similar
TITLE OF INVENTION: Intercellular Transactivating Strategies
NUMBER OF SEQUENCES: 38
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                                                                                                                                                                  Local Similarity
nes 8; Conserv
                                                                                                                                                                                                                                                                                                  TYPE: amino acid
TOPOLOGY: unknown
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STREET: Spring House Corpora
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72.7%;
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                                                                                                                                                                                  Score 49; DB 5;
Pred. No. 0.081;
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Pred. No. 0.081;
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          Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB seq length: 0
DB seq length: 2000000000
          protein search, using sw model
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Match
          PIR_71:*
1: pir1:*
2: pir2:*
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4: pir4:*
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Listing first 45 summaries
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144
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Copyright (c) 1993 - 2000 Compugen
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Ig kappa chain v r
Ig light chain V r
Ig kappa chain V r
Ig light chain V r
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Query Match
Best Local Similarity 100.0%; I
Matches 18; Conservative 0;
Y 12 SGSGTDFTLTISSLOAED 29

Score 87; DB Pred. No. 0. 0; Mismatches

DB 2; Le

Length 92, 1; 0; Indels

0

Gaps

0;

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RESULT 2 \$37530 STy530 If kappa chain V region (V-kappa 4) - human (fragment) C.Species: Homo sapiens (man) C.Species: Homo sapiens (man) C.Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999 C.Accession: \$37530 R.Klein, U.; Kueppers, R.; Rajewsky, K. submitted to the EMBL Data Library, September 1993 A.Description: Human IgM(+)IgD(+) cells, the major B cell subset in the peripheral A.Reference number: \$37501 A.Reference number: \$37501 A.Accession: \$3750 A.Status: preliminary A.Molecule type: mRNA A.Residues: 1-92 <kled a.cross="" c.keywords:="" c.superfamily:="" embl.226630;="" heterotetramer;="" homology="" immunoglobulin="" immunoglobulin<="" nid:9405702;="" pid:9405703="" pidn:caa81383.1;="" references:="" region;="" th="" v=""><th>Query Match 60.4%; Score 87; DB 2; Length 92; Best Local Similarity 100.0%; Fred. No. 0.00011; Matches 18; Conservative 0; Mismatches 0; Indels 0; Gaps 0; Qy 12 SGSGTDFTLTISSLQAED 29 1 </th><th>RESULT 1 S37533 Ig kappa chain V region (V-kappa 4) - human (fragment) C; Species: Homo sapiens (man) C; Species: Homo sapiens (man) C; Date: 06-Jan-1995 *sequence_revision 06-Jan-1995 *text_change 23-Jul-1999 C; Accession: S37533 R; Klein, U; Kueppers, R.; Rajewsky, K. submitted to the EMBL Data Library, September 1993 A; Description: Human IgM(+)IgD(+) cells, the major B cell subset in the peripheral A; Reference number: S37501 A; Reference number: S37501 A; Residues: 1-92 <kle> A; Molecule type: mRNA A; Residues: 1-92 <kle> A; Cross-references: EMBL: 226627; NID: 9405708; PIDN: CAA81380.1; PID: 9405709 C; Superfamily: immunoglobulin V region; immunoglobulin homology C; Keywords: heterotetramer; immunoglobulin</kle></kle></th><th>30 84 58.3 103 2 PH1051 31 84 58.3 103 2 PH1052 32 84 58.3 103 2 PH1054 33 84 58.3 103 2 PH1054 34 84 58.3 103 2 PH1047 35 84 58.3 104 2 PH1053 36 84 58.3 104 2 PH1102 37 84 58.3 104 2 PH1102 38 84 58.3 104 2 PH1103 39 84 58.3 104 2 PH1103 30 84 58.3 104 2 PH1104 31 84 58.3 104 2 PH1104 32 84 58.3 104 2 PH1104 33 84 58.3 104 2 PH1104 40 84 58.3 107 2 PH1104 41 84 58.3 107 2 PH1105 42 84 58.3 107 2 PH1105 43 84 58.3 107 2 PH1105 44 84 58.3 107 2 PH1105 45 86 58.3 107 2 PH1105 46 58.3 107 2 PH1105 47 88 58.3 107 2 PH1105 48 58.3 107 2 PH1105 49 58 58 58 58 58 58 58 58 58 58 58 58 58</th></kled>	Query Match 60.4%; Score 87; DB 2; Length 92; Best Local Similarity 100.0%; Fred. No. 0.00011; Matches 18; Conservative 0; Mismatches 0; Indels 0; Gaps 0; Qy 12 SGSGTDFTLTISSLQAED 29 1	RESULT 1 S37533 Ig kappa chain V region (V-kappa 4) - human (fragment) C; Species: Homo sapiens (man) C; Species: Homo sapiens (man) C; Date: 06-Jan-1995 *sequence_revision 06-Jan-1995 *text_change 23-Jul-1999 C; Accession: S37533 R; Klein, U; Kueppers, R.; Rajewsky, K. submitted to the EMBL Data Library, September 1993 A; Description: Human IgM(+)IgD(+) cells, the major B cell subset in the peripheral A; Reference number: S37501 A; Reference number: S37501 A; Residues: 1-92 <kle> A; Molecule type: mRNA A; Residues: 1-92 <kle> A; Cross-references: EMBL: 226627; NID: 9405708; PIDN: CAA81380.1; PID: 9405709 C; Superfamily: immunoglobulin V region; immunoglobulin homology C; Keywords: heterotetramer; immunoglobulin</kle></kle>	30 84 58.3 103 2 PH1051 31 84 58.3 103 2 PH1052 32 84 58.3 103 2 PH1054 33 84 58.3 103 2 PH1054 34 84 58.3 103 2 PH1047 35 84 58.3 104 2 PH1053 36 84 58.3 104 2 PH1102 37 84 58.3 104 2 PH1102 38 84 58.3 104 2 PH1103 39 84 58.3 104 2 PH1103 30 84 58.3 104 2 PH1104 31 84 58.3 104 2 PH1104 32 84 58.3 104 2 PH1104 33 84 58.3 104 2 PH1104 40 84 58.3 107 2 PH1104 41 84 58.3 107 2 PH1105 42 84 58.3 107 2 PH1105 43 84 58.3 107 2 PH1105 44 84 58.3 107 2 PH1105 45 86 58.3 107 2 PH1105 46 58.3 107 2 PH1105 47 88 58.3 107 2 PH1105 48 58.3 107 2 PH1105 49 58 58 58 58 58 58 58 58 58 58 58 58 58
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RESULT 3

S37529

Ig kappa chain V region (V-kappa 4) - human (fragment)

C; Species: Homo sapiens (man)

C; Species: Homo sapiens (man)

C; Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999

C; Accession: S37529

R; Klein, U.; Kueppers, R.; Rajewsky, K.

submitted to the EMBL Data Library, September 1993

A; Description: Human IgM(+)IgD(+) cells, the major B cell subset in the periphes A; Reference number: S37501

A; Reference number: S37501

A; Accession: S37529

A; Status: preliminary

A; Molecule type: mRNA

A; Residues: 1-92 KKLE>

A; Residues: 1-92 KKLE>

A; Residues: 1-92 KKLE>

A; Residues: heteroces: EMBL: 226631; NID: 9405700; PIDN: CAA81384.1; PID: 9405701

C; Reywords: heterotetramer; immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ig Kappa chain V region (V-kappa 4) - human (fragment)
C:Species: Homo sapiens (man)
C:Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
C:Accession: S37535
R:Klein, U.; Kueppers, R.; Rajewsky, K.
submitted to the EMBL Data Library, September 1993
A:Description: Human IgM(+)IgD(+) cells, the major B cell subset in the periphe A;Reference number: S37501
A;Reference number: S37501
A;Ccession: S37535
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-92 KLE>
A;Cross-references: EMBL: 226625; NID: g405712; PIDN:CAA81378.1; PID: g405713
C;Superfamily: Immunoglobulin V region; Immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
RESULT 5
37531
S37531
Ig kappa chain V region (V-kappa 4) - human (Iragment)
C:Species: Homo sapiens (man)
C:Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-C:Accession: 337531
R:Klein, U: Kueppers, R.: Rajewsky, K.
submitted to the EMBL Data Library, September 1993
A:Description: Human 1gM(+)1gD(+) cells, the major B cell subset in A:Reference number: 337501
A:Reference number: 337501
A:Status: preliminary
A:Molecule type: mRNA
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R:Klein, U.: Kueppers, R.: Rajewsky, K. submitted to the EMBL Data Library, September 1993 A;Description: Human IgM(+)IgD(+) cells, the major A;Reference number: S37501 A;Accession: S37534
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C;Date: 06-Jan-1995 *sequence_revision 06-Jan-1995 *text_change
C;Accession: S37534
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                                                                                                                                                                      A; Molecule type: mRNA
A; Residues; 1-92 KLEP
A; Cross-references: EMBL: Z26628; NID: g405706; PIDN: CAA81381.1; PID: g405707
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
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C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
C;Accession: S37532
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A;Molecule type: mRNA
A;Residues: 1-92 <KLE>
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A; Accession: S37532
A; Status: preliminary
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0.00011;
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0.00011;
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A49138
IqA kappa rheumatoid factor variable - human C; Species: Homo sapiens (man)
C; Date: 19-Dec-1993 #sequence_revision 18-Nc C; Paccession: A49138
R; Gause, A.; Kuppers, R.; Mierau, R.
Clin. Exp. Immunol. 88, 430-434, 1992
A; Title: A somatically mutated V kappa IV ge A; Reference number: A49138; MUID: 92298590
A; Accession: A49138
A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: protein
A;Residues: 1-102 (MIH)
C;Superfanily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
E;16-96/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B34153
If kappa chain V-IV region (Fue) - human
C;Species: Homo sapiens (man)
C;Date: 11-Apr-1990 *sequence_revision 11-Apr-1990 *text_change 21-Jan-2000
C;Accession: B34153
R;Mihaesco, E.; Ayadi, H.; Congy, N.; Gendron, M.C.; Roy, J.P.; Heyermann, H.; Frangione J. Biol. Chem. 264, 21481-21485, 1989
A;Title: Multiple mutations in the variable region of the kappa light chains of three mc A;Reference number: A34153; MUID:90094313
A;Accession: B34153
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C:Species: Homo sapiens (man)
C:Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 21-Jan-2000
C:Accession: PH0869
R:Manhelmer-Lory, A.; Katz, J.B.; Pillinger, M.; Ghossein, C.; Smith, A.; Diamond, B.
R:Title: Molecular characteristics of antibodies bearing an anti-DNA-associated idiotype A; Reference number: PH0862; MUID:92078875
A; Accession: PH0869
A; Reference number: PH0862; MUID:92078875
A; Accession: PH0869
A; Residues: 1-101 <AAN>
A; Residues: 1-101 <AAN>
A; Residues: 1-101 <AAN>
A; Rosidues: 1-101 <AAN>
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F;24-40/Region: complementarity-determining 1
F;41-55/Region: framework 2
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F:56-62/Region:
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Pred. No.
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Pred. No.
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A;Reference number: S3053
A;Accession: S30520
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-113 <MAR>
                                                                                                               A;Cross-references: EMBL;Z18325
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;16-96/Domain: immunoglobulin homology <IMM>
Ş
                                                                                                                                                                                                                                                                                                             Ig kappa chain V region - human c;Species: Homo sapiens (man) c;Date: 06-Jan-1995 #sequence_r C;Accession: S30520
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C;Superfamily: immunoglobulin V region; imm
C;Keywords: heterotetramer; immunoglobulin
F;16-96/Domain: immunoglobulin homology <IN
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A; Residues: 1-106 <GAU>
A; Residues: 1-206 <GAU>
A; Residues: 1-206 <GAU>
A; Cross-references: GB:S37926; NID:g298207; PIDN:AAB22366.1; PID:g298208
A; Note: sequence inconsistent with the nucleotide translation
A; Note: sequence extracted from NCBI backbone (NCBIN:106633, NCBIP:106637)
C; Superfamily: immunoglobulin V region; immunoglobulin homology
F; 20-99/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                            submitted to the EMBL Data Library, A; Reference number: S30520
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A;Molecule type: mRNA
A;Residues: 1-113 <MAR>
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C;Date: 06-Jan-1995 #sequence
C;Accession: S30523
R;Mariette, X.
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number: S30520
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                                                 Score 87;
Pred. No.
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pred. No. 0.00013;
0; Mismatches 0;
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Pred. No.
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71

SGSGTDFTLTISSLQAED

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RESULT 14

$34002

$34002

Ig kappa chain V region - human

C;Specles: Homo sapiens (man)

C;Date: 02-Dec-1993 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000

C;Accession: $34002; $30522

R;Mariette, X; Tsapis, A; Brouet, J.C.

Eur. J: Immunol. 23, 846-851, 1993

A;Title: Nucleotidic sequence analysis of the variable domains of four human monoclonal

A;Reference number: $34001; MUID:93209281

A;Accession: $34002

A;Accuse: preliminary

A;Molcoule type: mRMA

A;Residues: 1-113 <MAR>

A;Cross-references: EMBL:218328

C;Superfamily: immunoglobulin V region: immunoglobulin homology

C;Keywords: heterotetramer; immunoglobulin homology <IMM>
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$34003

If kappa chain V region - human
C; Species: Homo sapiens (man)
C; Accession: $34003
R; Mariette, X: Tsapis, A.; Brouet, J.C.
Eur. J. Immunol. 23, 846-851, 1993
Ritle: Nucleotidic sequence analysis of the variable domains of four human monoclonal
A; Reference number: $34001; MUID:93209281
A; Accession: $34003
A; Status: preliminary
A; Molecule type: manA
A; Residues: Lype: manA
A; Residues: 1-113 <MAR>
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 16-96/Domain: immunoglobulin homology <IMM>
KAHULN

C:Species: Homo sapiens (man)

C:Species: Homo sapiens (man)

C;Oate: 24-Apr-1984 #sequence_revision 24-Apr-1984 #text_change 01-Dec-2000

C;Accession: A01903; F61458

R;Schneider, M; Hilschmann, N.

Hoppe-Seyler's Z. Physiol. Chem. 356, 507-557, 1975

A;Title: Die Primaerstruktur einer monoklonalen Immunglobulin-i-Kette der Sta; Accession: A01903; MUID:76004342

A;Accession: A01903

A;Accession: A01903

A;Molecule type: protein
A;Note: this is the first completely sequenced V region of a new kappa chair
A;Note: this is the first completely sequenced V region of a new kappa chair
A;Note: this is the first completely sequenced V region of a new kappa chair
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Local Similarity 100.0%;
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Search completed: Job time: 469 sec

July 15,

2002, 13:01:03

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A; Molecule type: protein
A; Residues: 1-13 < SRO>
G; Comment: This is a Bence Jones protein.
G; Complex: An immunoglobulin heterotetramer subunit consists of two identical light (C; Complex: An immunoglobulin heterotetramer; Such as IgA and IgM, the subunits associate into the city of the consist of two identical light (C; Superfamily: immunoglobulin vegion; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin homology
C; Keywords: immunoglobulin homology < IMM>
C; Superfamily: Immunoglobulin homology < IMM>
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Copyright (c) 1993 - 2000 Compugen Ltd
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RESULT 2
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GSGTDF	:ch 11 Simi 18;	5 5	REVISION TO Salomon A.; Submitted (/ - - MISCELL/ - MISCELL/	SEQUENCE. SEQUENCE. MEDLINE=7600434 Schneider M., H "The primary st subgroup IV of Hoppe-Seyler's [2]	HUMAN 5; 1-1986 1-1996 1-1999 1-1999 1990 chas sapiens yota; 1 11a; Eu		73 72 71 71 71 70 70 70 70
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8 9	0	Benc	. > 5	5099 N.; a n type	Created) Last seq Last ann gion Len ordata; imates;		
	Sco Pr	Bence-Jones protein FRAMEMORK-1. COMPLEMENTARITY FRAMEMORK-2. COMPLEMENTARITY FRAMEWORK-3. COMPLEMENTARITY FRAMEWORK-4. BY SIMILARITY. MW: 0647F1D17F236	WISS-I	995; ; monoclonic pe (Bence-J Chem. 356:5	PRT; sed) sequence annotation Len. Len. cranial ss; Catarri	ALIGNMENTS	KY1X_HUMAN KY1L_HUMAN KY1L_HUMAN KY1R_HUMAN KY1R_HUMAN KY3B_HUMAN KY3B_HUMAN KY3B_HUMAN KY3B_HUMAN KY1T_HUMAN KY1T_HUMAN KY1T_HUMAN KY2B_HUMAN
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01-JAN-1988 (Rel. 06, Created)
01-JAN-1988 (Rel. 06, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
15 kappa chain V-IV region precursor (Fragment).
                                                                                            KV4B_HUMAN STANDARD; PRT; 13 AA.
P06313;
01-JAN-1988 (Rel. 06, Created)
01-JAN-1989 (Rel. 06, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
15-JUL-1999 (Rel. 38 Last annotation update)
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 SEQUENCE FROM N.A.
MEDLINE-86041853; PubMed-2997712;
Klobeck, H.G., Bornkamm G.W., Comb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
NCRI_TaxID-9606;
                                                                                   Homo sapiens (Human)
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InterPro; IPR003596;
Pfam; PF00047; ig; 1.
SMART; SM00406; IGV;
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PIR; A01902; K4HU.
HSSP; P01789; IMCP.
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MEDLINE-86041853; PubMed-2997712;
Klobeck H.G., Bornkamm G.W., Comb
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MISCELLANEOUS: THERE IS ONLY ONE IC KAPPA V-IV
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COMPLEMENTARITY-DETERMINING-3
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                                                             Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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Pfam; PF00047; Ig; 1.
SMART; SM00406; IGv; 1.
Immunoglobulin "
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P06314;
01-JAN-1988
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Nucleic Acids Res. 13:6515-6529(1985).
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Mammalia, Eutheria; Primates; Catarrhini; Hominidae,
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Ol-APR-1988 (Rel. 07, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
19 kappa chain V-TV region B17 precursor.
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                                                                           Submitted (OCT-1986)
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                                                                                                                                                                           SEQUENCE FROM N.A.
MEDLINE-86041854; PubMed-2997713;
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                                                                                        Marsh
                                                                                                     REVISION TO 76
                                                                                                                            Nucleic Acids Res.
                                                                                                                                                                                                                 NCBI_TaxID=9606;
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                                                                                                                                              F., Gould H.;
unique human V kappa IV
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Pred. No.
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COMPLEMENTARITY-DETERMINING-3.
FRAMEWORK-4.
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FRAMEWORK-2.
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Best Local S
Matches 18
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InterPro; IPR003596; 1
Pfam; PF00047; ig; 1
SMART; SM00406; IGv; 1
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"The primary structure of a monoclonal kappa-type chain of subgroup I (Bence-Jones Protein Hau): sul
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21-JUL-1986 (Rel. 01, Last
15-JUL-1999 (Rel. 38, Last
19 kappa chain V-I region
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P01600;
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PIR; A01905; K4HU17.
HSSP; P01789; 1MCP.
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SMART; SM00406; IGV;
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                                                                                                                                                                                                                                                                                     PIR; A01868; K1HUHU.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
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Pred. No.
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COMPLEMENTARITY-DETERMINING-1.
                                    FRAMEWORK-4
BY SIMILARI
                                                                   COMPLEMENTARITY - DETERMINING - 2. FRAMEWORK - 3. COMPLEMENTARITY - DETERMINING - 3.
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BY SIMILARI
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COMPLEMENTARITY-DETERMINING-2.
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bdivision within
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RESULT 7
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KV3J_HUMAN
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                                                                                                                                                                                                                                                                                                                                              EMBL; X02725; -; NOT_ANNOTATED_CDS.
PIR; A01901; K3HUVH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              904434;
13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
                                                                                                                                                             SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic Acids Res. 12:9229-9236(1984).
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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'; SM00406; IGv;
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COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
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Pred.
                                                                                                                   Score 82;
Pred. No.
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KV1I_HUMAN P01601;

STANDARD;

PRT;

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RESULT 8
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ID KVIW_H
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DT 13-AUG
DT 13-JUL
DE IG KAP
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Best Local Similarity 94.9
Matches 17; Conservative
13-AUG-1987 (Rel. C
13-AUG-1987 (Rel. O
15-JUL-1999 (Rel. 3)
Ig kappa chain V-T
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P04431;
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SEQUENCE
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EMBL; K01324; AAA58932.1; -.
EMBL; V00558; CAA73824.1; -.
PIR; A01881; K1HU11.
PIR; A21056; A21056.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the Buropean Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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MEDILINE-83129397; PubMed-6402305;
Bentley D.L., Rabbitts T.H.;
Bentley D.L., Rabbitts T.H.;
"Evolution of immunoglobulin V genes: evidence indicating Evolution of immunoglobulin V kappa sequences have diverged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
MEDLINE-6109856; PubMed-6779204;
Bentley D.L., Rabbitts T.H.;
"Human immunoglobulin variable region genes
                                                                                                                                                                                                                                                                                                                                             DOMAIN
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SIGNAL
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               kappa genes and a pseudogene.";
Nature 288:730-733(1980).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Mctazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Homo sapiens (Human)
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21-JUL-1986 (Rel.
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      05, Created)
05, Last sequence update)
38, Last annotation update)
I region Walker precursor.
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38, Last annotation update)
region HK101 precursor (Fragment).
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COMPLEMENTARITY-DETERMINING-1.
FRANEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
FRAMEWORK-3.
COMPLEMENTARITY-DETERMINING-3.
BY SIMILARITY.
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                                                                                                                                                                                            Score 82;
Pred. No.
                                                                     PRT;
                                                                                                                                                                                                                                            D7D0FF3718CEF587 CRC64
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2.7e-05;
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RESULT 9
KV3F_HUMAN
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Best Local
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P01624;
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Eukaryota; Metazoa; Chordata;
Mammalla; Eutheria; Primates;
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21-JUL-1986 (Rel. 01, Last sequence update)
16-CCT-2001 (Rel. 40, Last annotation updat
1g kappa chain V-III region POM.
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InterPro; IPR003596; Iç
Pfam; PF00047; ig; 1.
SMART; SM00406; IGV; 1.
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HSSP; P01607; 1REI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens (Human).
Eukaryota; Metazoa; C
Mammalia; Eutheria; F
                                                                                      SEQUENCE
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                                                                          MEDLINE=76276460; PubMed=60899;
                                                                                                          NCBI_TaxID=9606;
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Primates;
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Pred.
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COMPLEMENTARITY-DETERMINING-1.
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COMPLEMENTARITY-DETERMINING-2
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InterPro;

; IPR003006; ; IPR003596;

Ig_v. Ig_MHC HSSP; P01789; 2MCP

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RESULT 10

KV1A_MOUSE
ID KV1ASA
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SMART; SM00406; IGv; 1.
Immunoglobulin V region.
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SEQUENCE
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SMART; SM00406; IGv; 1.
Immunoglobulin V region
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InterPro; IPR003596; Ig_v.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       use by mon-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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MEDLINE=B1241357; PubMed=6788890;

Kwan S.-P., Rudikoff S., Seidman J.G., Leder P., Scharff M.D.;

"Nucleic acid and protein sequences of phosphocholine-binding light"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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Mus musculus (Mouse).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P01632;
21-JUL-1986 (Rel.
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16; Conserv
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88.9%;
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Pred. No. 5.2e
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Pred. No.
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COMPLEMENTARITY-DETERMINING-2.
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COMPLEMENTARITY-DETERMINING-1.
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5.2e-05;
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Query Match

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                                                                                                        DOMAIN
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Blochem. J. 171:337-347(1978).

-i- MISCELLANEOUS: THE MATURE CHAIN HAS 12 ADDITIONAL RESIDUES AT ITS
AMINO END, DUE TO A TANDEM DUPLICATION OF 36 NUCLEOTIDES AFTER THE
CODON FOR RESIDUE 36. RESIDUE 42 CORRESPONDS TO THE AMINO-TERMINAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  P01633;
21-JUL-1986
21-JUL-1986
15-JUL-1999
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DOMAIN
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MEDLINE-8 Colectough C., Perry
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                                               DOMAIN
                                                             DOMAIN
                                                                                               DOMAIN
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SMART; SM00406; IGV;
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InterPro; IPR003596; Ig_v.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Functional significance and evolutionary development of the 5'-terminal regions of immunoglobulin variable-region genes."; cell 29:681-689(1982).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ig kappa chain V-V region MPC11 precursor.
Mus musculus (Mouse)
 SEQUENCE
             NON_TER
                          REPEAT
                                    REPEAT
                                                                                                                                                                                                                  PIR; A01916; KVMS11.
                                                                                                                                                                                                                                                                                                                                                                                                                      "Sequence of the full-length immunoglobulin kappa-chain
                                                                                                                                                                                                                                                                                                                                                                                                                                      Smith G.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=78186617; PubMed=418775;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE OF 30-149.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "The variability, arrangement, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=10090;
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                                                                                                                                                       [mmunoglobulin V region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               71
                                                                                                                                                                                                                                                                                                                                                    RESIDUE OF TYPICAL KAPPA CHAINS.
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                                                                                                                                                       Signal; Repeat.
 ž
                                                         FRAMEWORK-3.
COMPLEMENTARITY-DETERMINING-3
                                                                              FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
                                                                                                        FRAMEWORK-1.
COMPLEMENTARITY-DETERMINING-1.
                                                FRAMEWORK - 4.
                                                                                                                                IG KAPPA CHAIN V-V REGION MPC11.
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B0480C87B682AC3E CRC64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R.P.;
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RESULT 12

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NON_TER
SEQUENCE
                                                                                                                   P04433:
13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation updat
15 kappa chain V-III region VG precursor (F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PIR; A01875; K1HUSW.
HSSP; P01607; 1REI.
InterPro; IPR003006;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             kappa-type (Bence Jones protein Scw.), and the complete amino acid sequence Hoppe-Seyler's Z. Physiol. Chem. 355.86-1- MISCELLANEOUS: THE C REGION OF THIS
                                                            Homo sapiens (Human).
Eukaryota; Metazoa; C
Mammalia; Eutheria; F
                                                                                                                                                                                                                           KV3I_HUMAN
                                                                                                                                                                                                                                               HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00047; 1g; 1.
SMART; SM00406; IGV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eulitz M., Hilschmann N.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ig kappa chain V-I region Scw. Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence up
15-JUL-1999 (Rel. 38, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P01609;
    SEQUENCE FROM N.A
                                         NCBI_TaxID-9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KV1Q_HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -1- MISCELLANEOUS: THIS IS A BENCE-JONES PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunoglobulin V region;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  106
                                                                                                                                                                                                                                                                                                                             65
                                                                                                                                                                                                                                                                                                                                                 12 SGSGTDFTLTISSLQAED 29
                                                                                                                                                                                                                                                                                                                                                                                                                               Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MISCELLANEOUS: THE C REGION OF THIS CHAIN HAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SGSGTDFTLTISSLQAED
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16; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      structure of a human immunoglobulin L-chain of Bence-Jones protein Scw.), II: The chymotryptic
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                                                                 Primates;
                                                                                Chordata;
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88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bence-Jones protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MW;
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Pred. No.
1; Mismatc
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1; Mismatches
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FRAMEWORK-3.
COMPLEMENTARITY-DETERMINING-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BY SIMILARITY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPLEMENTARITY-DETERMINING-1. FRAMEWORK-2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                          Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                             PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
                                                                                                                       precursor (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             32CECDDDF9644414 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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RESULT 14
KV1M_HUMAN
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DOMAIN
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InterPro; IPR003596; Ig_v.
pfam; PP00047; ig; 1.
SMARY; SW00406; IGv; 1.
Immunoglobulin V region; Sig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pech
                                                                                                                                           Scand. J. Immunol. 5:677-684 (1976).
-I- MISCELLANEOUS: THE SECOND AND THIRD HYPERVARIABLE REGIONS OF THIS CHAIN ARE IDENTICAL WITH THOSE OF THE HUMAN POM V-III KAPPA CHAIN.
                                                                                                                                                                                                         "Complete amino acid sequence of the variable of IgM anti-gamma globulins (Lay/Pom) with shared specificities.";
                                                                                                                                                                                                                                                                           Capra J.D.,
                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                         Ig kappa chain V-I region Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; X01668; -; NOT_ANNOTATED_CDS. PIR; A01900; K3HUVG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entities requires a license agreement (See http://www.isb-sib.ch/announce/
or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the European Bioinformatics Institute. The second of the s
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic Acids Res. 12:9229-9236(1984).
InterPro;
                                         HSSP;
                                                                                                  WITH WHICH IT SHARES CERTAIN IDIOTYPIC -i- MISCELLANEOUS: THIS CHAIN WAS ISOLATED
                                                                                                                                                                                                                                                                                                MEDLINE=77038198; PubMed=824717;
                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P01605;
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Immunoglobulin genes of different subgroups
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                  GLOBULIN ACTIVITY.
R; A01871; K1HULY.
SP; P01607; 1REI.
terPro; IPR003006; Ig
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16;
  IPR003596;
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                                                                                                                                                                                                                                                                         Klapper D.G.;
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Kratzin H., Yang C., Krusche J.U., Hilschmann N.;
"Preparative separation of the tryptic hydrolysate of a protein
high-pressure liquid chromatography. The primary structure of a
monoclonal L-chain of k-type, subgroup I (Bence-Jones Protein
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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; SM00406;
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1 23 FRAMEWORK-1.

24 34 COMPLEMENTARITY-DETERMINING-1.

35 49 FRAMEWORK-2.

50 56 COMPLEMENTARITY-DETERMINING-2.

57 88 FRAMEWORK-3.

89 97 COMPLEMENTARITY-DETERMINING-3.

98 107 FRAMEWORK-4.

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COMPLEMENTARITY - DETERMINING - 2.
FRAMEWORK - 3.
COMPLEMENTARITY - DETERMINING - 3.
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COMPLEMENTARITY-DETERMINING-1.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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69	69	70	70	70	71	73	74	77	79	08	08	82	82	82	83	Score	
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MEDITINE-98277139; PubMed=9614934;
MEDITINE B., Van der Merwe P.L.,
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Clin. Immunol. Immunopathol. 87:184-192(1998).
EMBL: AF035033; AAD56209.1; -.
HSSP; P01607; 1REI.
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OSUL81; PREVIDENTAL: 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Cast sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
(FRAGMENT).
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Pfam; PF00047; ig; 1.
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Pred. No. 3.2e-05;
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Best Local
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Q9UL77;
01-MAY-2000 (TrEMBLrel. 1
01-MAY-2000 (TrEMBLrel. 1
01-DEC-2001 (TrEMBLrel. 1
01-DEC-2001 (TrEMBLrel. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fetus.";
Clin. Im
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SEQUENCE FROM N.A.
MEDLINE-98277139; PubMed-9614934;
MEDLINE-98277139; PubMed-9614934;
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01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last snnotation update)
01-DEC-2001 (TrEMBLrel. 19, Last snnotation update)
ANTI-STREPTOCOCCAL/ANTI-MYOSIN IMMUNOGLOBULIN KAPPA
VARIABLE REGION (FRAMENT).
Homo sapiens (Human)
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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Adderson B.E., Shikhman A.R., ward K.E., Cunningham M.W.;
"Molecular analysis of polyreactive monoclonal antibodies from rheumatic carditis: human anti-N-acetylglucosamine/anti-myosin antibody v region genes.";
J. Immunol. 161:2020-2021(1998).
EMBL; U96396; AAB68785.1; ".
                                                                                                                                                                                                                                                                                                                 Pfam; PF00047; 1g; 1
SMART; SM00406; IGv;
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HSSP; P01607; 1REI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wu X., Llu B., Van Young D.C.;
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Mammalla; Eutheria;
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                                                                                                                                                                                                     SEQUENCE
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InterPro; JPR003596;
Local Similarity
nes 17, Conser
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107 AA;
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56.98;
ilarity 94.48;
Conservative
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EMBLrel. 13, Last sequence update)
EMBLrel. 19, Last annotation update)
IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
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11520 MW;
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Primates;
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Primates;
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Score 82; DB
Pred. No. 4.6e
0; Mismatches
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Catarrhini; Hominidae; Homo.
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ed. No. 4.56
Mismatches
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4.5e-05;
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                                                                                                                            "Cloning genes of J. Cell.
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Q9UL70;
Q1-MAY-2000 (TrEMBLrel. 13, Created)
Q1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
Q1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
                                                                                                                                                                                SEQUENCE
Chen P.,
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Q9ERZ9;
01-MAR-2001 (TIEMBLITE1. 16, Created)
01-MAR-2001 (TIEMBLITE1. 16, Last sequence update)
01-DEC-2001 (TIEMBLITE1. 19, Last annotation update)
ANTI HUMAN TNF-ALDHA LIGHT CHAIN VARIABLE REGION (FI
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Mammalia; Eutheria;
NCBI_TaxID=10090;
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SMART; SM00406; IGv; 1.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae
                                              SEQUENCE FROM N.A.

Chen P., Deng J.B., Wang Z.L., Han H., Yao L.B.,

"Construction and sequencing of the single-chain

human TNF-alpha specific monoclonal antibody.";
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SEQUENCE FROM N.A
                                                                                                                                                                                                                                                                                 Mus musculus (Mouse).
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InterPro; IPR003596; Ig_v.
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                                                                                                                                                                              Deng J.B.,
                                                                                                                            Deng J.B., Wang Z.L., Han H., Su C.Z.;
and sequencing of the light chain fragment
an anti-hTMP-a monoclonal antibody.";
Mol. Immunol. 12:21-26(1996).
                                i Ta Hsueh
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108 AA;
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Sciurognathi; Muridae; Murinae; Mus
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SMART; SM00409; IG; 1.
SMART; SM00406; IGv; 1.
NON_TER 107 107
SEQUENCE 107 AA; 1179
                                   Q9UL79
Q9UL79;
Q9UL79;
Q1-MAY-2000
01-MAY-2000
01-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                              NON_TER
NON_TER
SEQUENCE
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SEQUENCE FROM N.A.
MEDLINE-98277139; PubMed-9614934;
"''''' B., Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q9UL85;
Q1-MAY-2000 (TrEMBLrel. 13, Created)
Q1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
Q1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN KAPPA CHAIN VARIABLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro;
InterPro;
MYOSIN-REACTIVE (FRAGMENT).
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SMART; SM00406; IGv; 1.
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InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HSSP; P80362; 1WTL.
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IPR003006; Ig_MHC.
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IPR003596;
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109
             (TYEMBLrel. 13, Created)
(TYEMBLrel. 13, Last sequence update)
(TYEMBLrel. 19, Last annotation update)
TIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                        PRELIMINARY;
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11761 MW;
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Pred. No. 9.4e
2; Mismatches
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                                                                                                                        PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT;
                                                                                                                                                                                                                                                                                                                                                                                                              FB1E43E7C7AFACCC CRC64;
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No. 9.4e-05;
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Best Local S
Matches 16
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SEQUENCE
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01-MAY-2000 (TrEMBLrel. 13, Created)
01-DEC-2001 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation updat
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIAB
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MEDLINE=98277139; PubMed=9614934;
MEDLINE=98277139; PubMed=9614934;
MEDLINE=98277139; PubMed=9614934;
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SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934;
DL. Tin B., Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Clin. Immunol. Immunol. EMBL; AF035035; AAI HSSP; P01607; 1REI
                                                                                                                     InterPro; IPA003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
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                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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SMART; SM00406; IGV; 1.
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                                                                                                                                                                                             fetus.";
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                                                                                                                                                                                                                                                                   NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                             (FRAGMENT).
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           SGSGTDFTLTISSLQAED
SGSGTEFTLTISSLQFED
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                                         l Similarity
16; Conser
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108 AA;
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108 AA;
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                                                                                                                                                                        Immunopathol.
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11834 MW;
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11787
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                                                  53.5%;
88.9%;
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Pred. No.
                                                 Score 77; DB 4;
Pred. No. 0.00027;
                                                                                                                                                                                   87:184-192(1998)
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RESULT OF REAL PROPERTY OF THE PROPERTY OF THE
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Matches 15
   Query Match
                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

MEDILNE-20250927; PubMed-10788485;

Rader C., Ritter G., Nathan S., Elia M., Gout
Cohen L.S., Welt S., Old L.J., Barbas C.F. II

"The rabbit antibody repertoire as a novel so
of therapeutic human antibodies.";
J. Biol. Chem. 275:13668-13676(2000).

EMBL; AF245502; AAF68449.1;
HSSP: P8036; IWTL.
HSSP: P80362; IWTL.
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Q9UL78;
Q1-MAY-2000 (TrEMBLrel. 13, Created)
Q1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
Q1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOSLOBULIN LIGHT CHAIN VARIABLE
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01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
ANTI-HUMAN A33 LIGHT CHAIN VARIABLE REGION (FRAGMENT).
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oryctolagus cuniculus (Rabbit)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
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                                                                                                                                                                                        SMART;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9986;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR003006;
                                                                                                                                                                                                               InterPro; IPR003596; Pfam; PF00047; 19; 1
                                                                                                                                                                                                                                                                               InterPro; IPR003006;
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                                                                                                                                                                                     SM00406;
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15; Conserv
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00047; 1g; 1.
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109 AA;
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                                                                                          3
                                                                                                                                                                                     IGV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               109
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                                                                                             11323
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           51.4%;
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   . 78
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                                                                                             WW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 74; DB 4;
Pred. No. 0.0008;
1; Mismatches
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   Score
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Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
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                                                                                                 BD8B396EE75F94FB
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   73;
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   DВ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gout I.,
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RESULT
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                                                                                                                                                                                                                                                                                                                                                       09JL82;
01-0CT-2000
01-0CT-2000
01-DEC-2001
            STRAIN-BALB/C;
MEDLINE-20448942; PubMed-10992488;
Malkiel S., Liao L., Cunningham M.W., Diamond B.;
"T-Cell-dependent antibody response to the dominant streptococcal polysaccharide, N-acetyl-glucosamine, with cardiac myosin.";
Infect: Immun. 68:5803-5808(2000).
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Mus musculus (Mouse).

Mus musculus (Mouse).

Charyota; Metazoa; Chordata; (
                                                                                                                                                                                                                  Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10090;
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MalkLel S., Liao L., Conningham M.W., Diamond B.;
"I-cell-dependent antibody response to the dominant
streptococcal polysaccharide, N-acetyl-glucosamine,
with cardiac myosin.";
Infect I
                                                                                                                                                                                                                                                                                                                                 01-0CT-2000 (TrEMBLrel. 15, 01-0CT-2000 (TrEMBLrel. 15, 01-DEC-2001 (TrEMBLrel. 19, ANTI-MYOSIN IMMUNOGLOBULIN )
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SMART; SM00406; IGV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Infect. Immun. 68:5803-5808(2000).
EMBL; AF206032; AAF69330.1;
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
ANTI-MYOSIN IMMUNOGLOBULIN LIGHT CHAIN VARIABLE REGION
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01-OCT-2000
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                                                                                                                                                                                                                                                                                                                (FRAGMENT)
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  AF206024;
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LIGHT CHAIN VARIABLE REGION
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Pred.
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RESULT
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DT QQ
DT QQ
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RESULT
ID Q99M37 PRELIMINARY; PRT; 238 AA.
AC Q99M37; Q99M37; PRT 238 AA.
AC Q99M37; PRT 2001 (TrEMBLrel. 17, Created)
DT 01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
DT 01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
DE HYPOTHETICAL 26.3 KDA PROTEIN.
OS Mus musculus (Mouse).
C Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
C Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
X NCBL_TaxID=10090;
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Best Local S
Matches 14
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Best Local Similarity
Matches 14; Conserv
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InterPro; IPR003596; Ig_N
Pfam; PF00047; Ig; 1.
SMART; SM00406; IGV; 1.
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InterPro; IPR003006; Ig
InterPro; IPR003596; Ig
InterPro; IPR003596; Ig
Pfam; PF00047; 1g; 1
SMART; SM00406; IGv; 1
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01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MYOSIN-REACTIVE IMMUNOGLOBULIN LIGHT CHAIN VARIABLE
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MEDILINE-98277139; PubMed-9614934;
Mu X., Liu B., Van der Merwe P.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NON_TER
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (Human)
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.: 12775 MW;
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Primates;
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77.8%;
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Pred. No. 0.00
2; Mismatches
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        070E31E210D1CB01 CRC64;
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Best Local S
Matches 13
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Best Local S
Matches 14
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InterPro; IPR003597; Ig_cl.
InterPro; IPR003500; Ig_like.
InterPro; IPR003506; Ig_MHC.
InterPro; IPR003006; Ig_v.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; Ig; 2
SMART; SM00409; IG; 2
SMART; SM00407; IGC1; 1.
SMART; SM00401; IGC; 1.
SMART; SM00401; IGC; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q91WS9 PRELIMINARY; PRT; 233 & Q91WS9; O1-DEC-2001 (TrEMBLrel. 19, Created) O1-DEC-2001 (TrEMBLrel. 19, Last sequence O1-DEC-2001 (TrEMBLrel. 19, Last annotatic HYPOTHETICAL 25.8 KDA PROTEIN (FRAGMENT).

Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
NCBI_TaxID-10090;
                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (SEP-2001) to the EMBL; BC013496; AAH13496.1;
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Hypothetical protein.
SEQUENCE 238 AA; 26344 MW; FB2BC
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NON_TER 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM
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completed: July
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                                                                                                                                                                                                                ch 47.9%;
l Similarity 72.2%;
l3; Conservative
                                                                                                                                                                                                                                                                                                                                           233 AA;
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                       15,
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77.8%;
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Last annotation updat
                                                                                                                                                                                                                Score 69; DB
Pred. No. 0.01
4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL/GenBank/DDBJ databases
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 2000000000
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1: \SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1980.DAT:*

2: \SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1981.DAT:*

3: \SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1982.DAT:*

4: \SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1982.DAT:*

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5: \SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1985.DAT:*

5: \SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1986.DAT:*

7: \SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1987.DAT:*

8: \SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1989.DAT:*

9: \SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1999.DAT:*

10: \SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1999.DAT:*

11: \SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1999.DAT:*

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13: \SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1993.DAT:*

14: \SIDS1/gcgdata/hold-geneseq/geneseqp-emb1/AA1993.DAT:*

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14.104 Million cell updates/sec
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Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2000 Com
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					SUMMAKIES	
		OMD.				
No.	Score	Query Match	Query Match Length	DB	ID	Description
1	87	60.4	82	19	AAW62807	Amino acid seque
2	87	60.4	94	19	AAW62808	
ω	87	60.4	107	20	AAY05269	
4	87	60.4	111	15	AAR52059	Light chain vari
S	87	60.4	112	18	AAW27527	
6	87	60.4	113	15	AAR50313	ised 1
7	87	60.4	113	15	AAR50314	_
8	87	60.4	113	17	AAR98489	Humanised 2B6 an
9	87	60.4	113	19	AAW42466	Interleukin-5 hu
10	87	60.4	113	21	AAB18857	Amino acid seque
::	87	60.4	113	21	AAB18861	Amino acid seque

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12
87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87	87
60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4	60.4
149	147	137	137	137	135	134	133	133	133	132	131	131	131	131	131	131	127	126	126	124	120	120	115	114	114	114	113	113	113	113	113		113
20	20	18	18	18	21	20	20	18	9	21	20	20	20	20	16	16	17	22	21	20	22	15	18	18	5	14	22	21	21	21	21	21	21
AAY34311	AAY34314	AAW10545	AAW10544	AAW10547	AAB03714	AAY50690	AAY05264	AAW21934	AAP80894	AAB07966	AAY18118	AAY18126	AAY23771	AAY23779	AAR70202	AAR75355	AAW06443	AAB78867	AAY77598	AAY57181	AAG65565	AAR61240	AAW27546	AAW15537	AAR45605	AAR30144	AAB62769	AAB18881	AAB18879	AAB18873	AAB18871	AAB18869	AAB18863
IgM antibody CEM 1	ody CEM	murine	murine	Wild type murine a	Immunoglobulin kap	4 VI 0	Antibody 24-31 hum	11g1		۳-	nain	chain	chain	-		antib	VL region.	human Flt-1					light c	Anti-TGF beta-1 sc	oclonal antibo	AH vari	HIV-1	acid	Amino acid sequenc			id	Amino acid sequenc

ALIGNMENTS

AAW62807

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AAW62807 standard; Peptide;

82 A

23-SEP-1998 (first entry)

AAW62807;

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Green L,
Mendez M;
                                                                                                                                                                                          Human; immunoglobulin; Ig; transgenic; non-human mammal; inactivated endogenous Ig locus; B-cell development; human heavy chain Ig locus; micro constant region; J-H; D-H; V-H gene; kappa light chain Ig locus; kappa constant region; J-kappa gene; V-kappa; production; antibody.
                                                                                          03-DEC-1996;
                                                                                                              03-DEC-1997;
New transgenic non-human mammals - having an inactivated
                     WPI; 1998-333314/29
                                                                       (ABGE-) ABGENIX INC.
                                                                                                                                  11-JUN-1998
                                                                                                                                                      WO9824893-A2
                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                         Amino acid sequence of a human antibody fragment.
                                                Jakobovits
                                                                                          96US-0759620
                                                                                                              97WO-US23091.
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                                                Klapholz
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                       Green L,
                 New transgenic non-human mammals - having an inactivated immunoglobulin locus and a near complete human immunoglo
                                                                                                                                                                                                                                                                                                                                                                                                      Human; immunoglobulin; Ig; transgenic; non-human mammal; inactivated endogenous Ig locus; B-cell development; D-H; D-H; V-H gene; human heavy chain Ig locus; micro constant region; J-K; D-H; V-H gene; kappa light chain Ig locus; kappa constant region; J-kappa gene; V-kappa;
 used
                                                                         WPI; 1998-333314/29
                                                                                                                                                                                                                                         03-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW62808 standard; Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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                                                                                                                                                                                                       03-DEC-1996;
                                                                                                                                                                                                                                                                               11-JUN-1998
                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                          production; antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Amino acid sequence of a human antibody fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 78; 128pp; English.
                                                                                                                                                                 (ABGE-) ABGENIX INC
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hes 18; Conser
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 tor
production of human antibodies
                                                                                                                             Jakobovits A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                             Klapholz S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94
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Pred. No.
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                                                                                                                             Kucherlapati
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. 5.6e-05;
                 human immunoglobulin locus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 82;
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Best Local s
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   locus, so that the mammal does not display normal B-cell development. The modified genome also has an inserted human heavy chain Ig locus in germline configuration, the human heavy chain Ig locus comprising a human micro constant region and regulatory and switch sequences, human J-H genes, human D-H genes, and human V-H genes and an inserted human kappa light chain Ig locus in germline configuration, the human kappa 19ht chain Ig locus in germline configuration, the human kappa 19ht chain Ig locus comprising a human kappa constant region, J-kappa genes, and V-kappa genes, where the number of V-H and V-kappa genes inserted are selected to restore normal B-cell development in the mammal. The transgenic animals have a near complete human Ig locus, including both a human heavy chain locus and a human kappa light chain locus. They can be used for the production of human antibodies when exposed to particular antigens e.g. when exposed to human II-6, EGFR or TNF- alpha to the mice will produce antibodies to II-8, EGFR or TNF- alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  transgenic Xenomice, created using the member of the transgenic non-human mammal which has genome specification describes a transgenic non-human mammal which has genome modifications that comprise an inactivated endogenous immunoglobulin (Ig) modifications that comprise an inactivated endogenous immunoglobulin (Ig) modifications that the mammal does not display normal B-cell development. The
Claim 7; Page 84; 122pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antibody 24-31; variable light chain; humanised antibody; gene therapy; CD40 binding inhibitor; gp39; autoimmune disease; food-related allergy; graff-versus-bost disease; graft rejection; intestinal inflammation; reversible obstructive airways disease; allergy; transplant rejection;
                                       New humanized antibodies to human gp30
                                                                                                                  Black A,
                                                                                                                                                                                                                                    08-SEP-1998;
                                                                                                                                                                                                                                                                                                             WO9912566-A1
                                                                                                                                                                                                                                                                                                                                                                                                                           immune response inhibitor; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antibody 24-31 humanised VL chain sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW62793-822 represent fragments of human antibodies produced by transgenic Xenomice, created using the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 79; 128pp; English
                                                                            WPI; 1999-229142/19
                                                                                                                                                       (IDEC-) IDEC PHARM CORP.
                                                                                                                                                                                             08-SEP-1997;
                                                                                                                                                                                                                                                                        18-MAR-1999
                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                        Mus sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                  asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY05269 standard; Protein; 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  leukaemia;
                                                                                                                  Hanna N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                             97us-0925339
                                                                                                                                                                                                                                    98WO-US18163
                                                                                                                  Newman RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                            lymphoma; immunosuppressant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   60.4%; Score 87; DB 100.0%; Pred. No. 6. Live 0; Mismatches
                                                                                                                  Padlan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 19; 1
. 6.4e-05;
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 thrombocytopenic purpura (ITP) or non-autoimmune conditions such as graft-versus host disease (GVHD) or graft rejection. They can also be used for the treatment of reversible obstructive airways disease, intestinal inflammations and allergies (e.g. coeliac disease, Crohn's disease and ulcerative colitis) and food-related allergies (e.g. migraine, rhinitis and eczema), transplant rejection, asthma, leukaemia, intermediate of the condition of the
  09-SEP-1992
                                                     07-SEP-1993;
                                                                                                                                                                                                                                                                                      Region
                                                                                                                                                                                                                                                                                                                                         Region
                                                                                                                                                                                                                                                                                                                                                                                         Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antibody; humanised; murine; human; heavy chain; light; variable; framework region; complementarity determining region; reshaping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAR52059 standard; Protein; 111 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or lymphoma. The antibodies can also be used as immunosuppressants, particular during gene or cellular therapy. They may be used to inh humoral and cellular immune responses against viral vectors, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence is the variable light chain of a humanised antibody of the invention. The antibody is capable of competing with a murine 24-31 antibody for inhibiting CD40 binding to gp39. The humanised antibodies can be used for the treatment of a disease treatable by modulating gp39 expression or inhibiting the gp39/CD40 interaction. They can be used for treating e.g. autoimmune diseases such as rheumatoid arthritis, psoriasis multiple sclerosis, diabetes, systemic lupus crythematosus and idiopathic
                                                                                                                                                        EP592106-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modelling; surface
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Light chain variable region of human KV4B antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 retroviral vectors or adenoviral vectors. The use of such antibodies should enable such cells or vectors to be administered repeatedly, which will facilitate treatment of chronic diseases such as cancers and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   autoimmune diseases.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
  92US-0942245
                                                  93EP-0307051.
                                                                                                                                                                                                                                                                                      /note-
63..95
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56..62
                                                                                                                                                                                                          /note-
                                                                                                                                                                                                                                                             /note=
                                                                                                                                                                                                                                                                                                                                                                                                                /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                   /label= complementarity_determining_region_1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
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                                                                                                                                                                                                          "CDR 3"
                                                                                                                                                                                                                                                             "FR 3"
                                                                                                                                                                                                                                                                                                          *CDR 2*
                                                                                                                                                                                                                                                                                                                                                                "FR 2"
                                                                                                                                                                                                                                                                                                                                                                                                                  "CDR 1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "FR 1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 107;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            to inhibit
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AAW27527
ID AAW
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                             (ASAK ) ASAHI
(NIKK-) NIKKA
(TORI ) TORII
(TSUR/) TSURA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Modification of a rodent antibody (Ab) or fragment by resurfacing in order to produce a humanised rodent Ab can be determined by calculating homology between murine and human Ab antibody surfaces. In order to test the resurfacing approach of the invention, three humanisation experiments were set up: (1) traditional loop grafting; (2) resurfacing approach using most similar chain; and (3) resurfacing approach using human sequences with most similar surface residues. The Ab used was the murine anti-N901 Ab (see AAR52055). Experiment 3 was carried out using the present sequence which represents the human KV4B Ab light chain variable region with 71 percent homology with anti-N901 Ab. N901/KV4B
      Humanised, semi-chimeric and chimeric antibodies against human high-affinity IqE receptor - useful medicinally and have low
                                                                                                                                                                                                                                                                                                 Complementarity determining region; CDR; murine; mouse; human; high affinity; immunoglobulin E; receptor; monoclonal antibody; IgE; MAb; light chain; variable region; humanised; semi-chimeri
                                                   WPI; 1997-429186/40.
                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                           chimeric; treatment;
                                                                                                                                                                                                                                                                                                                                                              Light chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Method of resurfacing of rodent antibodies to produce humanised antibody forms \cdot for producing non-human antibodies with improved therapeutic efficiency by presenting human surface on V-region
                                                                                                                                                    19-JAN-1996;
                                                                                                                                                                              19-JAN-1996;
                                                                                                                                                                                                         29-JUL-1997
                                                                                                                                                                                                                                    JP09191886-A
                                                                                                                                                                                                                                                                                                                                                                                           16-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                                      AAW27527;
                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW27527 standard; Protein; 112 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Fig 4A; 230pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1994-120230/15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Guild BC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (PEDE/) PEDERSEN J T. (IMMU-) IMMUNOGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (AAR52060)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  70 sgsgtdftltisslqaed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12 SGSGTDFTLTISSLQAED 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ഗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         was prepd. by resurfacing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pedersen JT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               111 AA;
                                                                                                                                                                                                                                                                                                                                                            variable region of human CRA2 antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                              YAKUHIN
T.
                                                                                                          WHISKEY
                                                                                                                       BREWERIES LTD
                                                                                                                                                   96JP-0024816
                                                                                                                                                                              96JP-0024816
                                                                                                                                                                                                                                                                                  prevention; disease; allergy; CRA2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       60.4%;
100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rees
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 15;
7.7e-05;
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high-affinity IgE receptor antigenicity in humans

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Best Local Similarity
Matches 18; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence, the light chain variable region of the human antibody (Ab) CRA2, was used in the preparation of a humanised or semi-chimeric monoclonal Ab (MAb), comprising complementarity determining regions (CDR) from a murine, anti-human high affinity immunoglobulin E (IgE) receptor, MAb. The humanised, semi-chimeric or chimeric MAb can be used to treat or prevent diseases, specifically allergies, associated with the receptor, and has very
                                                                                                                                                                                                                                                                                                                                                               Monoclonal antibody; Plasmodium falciparum; CDR; complementarity determining region; fusion protein; murine; variable; light; heavy; chain; malaria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
                     Claim 5;
                                                                                                             Charoenvit Y,
Sadoff JC, Sy
                                                                                                                                                     (USSA)
                                                                                                                                                                                   09-SEP-1992;
                                                                                                                                                                                                     08-SEP-1993;
                                                                                                                                                                                                                         17-MAR-1994
                                                                                                                                                                                                                                             W09405690-A
                                                                                                                                                                                                                                                                         Region
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                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                             05-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR50313;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AARS0313 standard; Protein; 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
 Naturally-occurring (AAQ44841-42) and synthetic (AAQ44825-28)
                                                  New engineered antibodies and fusion proteins plasmodium infection - contg. murine antibody
                                                                                N-PSDB; AAQ44827.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              low antigenicity in humans.
                                                                                          WPI; 1994-101115/12
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                                                                                                                                         SMITHKLINE BEECHAM CORP
US SEC OF ARMY.
US SEC OF NAVY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 14; 26pp; Japanese
                                        nucleic acid, vectors and transformed
                    Fig 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             112
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                                                                                                              Sylvester
                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                              Hoffman S,
lvester DR,
                                                                                                                                                                                   92US-0941654
                                                                                                                                                                                                      93WO-US08435
                  98pp;
                                                                                                                                                                                                                                                                          /label- CDR2
95..103
                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                /label= CDR3
                                                                                                                                                                                                                                                                                                       /label - CDR1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                       variable region Pfhzlc1-1
                     English
                                                                                                              Hurle
Gross
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ₽
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                                                                                                             MS,
                                                                                                                        Rosenberg
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7.8e-05;
                                                    CDR
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R sequences,
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RESULT
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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  complementarity determining region; fusion prot murine; variable; light; heavy; chain; malaria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
Naturally-occurring (AAQ44841-42) and synthetic (AAQ44825-28) variable light chain and variable heavy chain sequences derived from murine mAb NFS2 are provided. Murine mAb NFS2, its variable chain peptides, CDRs, functional fragments, Fab fragments, and analogs are useful in prodn. of fusion proteins, esp. engineered antibodies. These prods. are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR50314;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              used to protect humans against Plasmodium infections
                                                                                                    New engineered antibodies and fusion proteins for preventing plasmodium infection - contg. murine antibody CDR sequences, corresp. nucleic acid, vectors and transformed cells
                                                                                                                                                                                Charoenvit Y, Sadoff JC, S
                                                                                                                                                                                                                (SMIK ) SMITHKLINE BEECHAM (USSA ) US SEC OF ARMY. (USNA ) US SEC OF NAVY.
                                                                                                                                                                                                                                                               09-SEP-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-OCT-1994
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                                                                             Claim
                                                                                                                                                N-PSDB; AAQ44828.
                                                                                                                                                                                                                                                                                                                                                                                                          Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Monoclonal antibody; Plasmodium falciparum; CDR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Humanised light chain variable
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                                                                                                                                                           1994-101115/12
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                                                                             Fig
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                                                                                                                                                                                 Sylvester
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                                                                                                                                                                                            Hoffman
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95..103
/label= CDR3
                                                                                                                                                                                                                                                                                                                                                                                     /label-
56..62
                                                                              98pp; English
                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers 24..40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protein;
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100.08;
                                                                                                                                                                                 DR,
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                                                                                                                                                                                 Hurle
Gross
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               fusion protein;
                                                                                                                                                                                               Rosenberg
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7.8e-05;
hes 0;
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                                                         Ames RS, ...
                                                                                                          06-JUN-1995;
23-DEC-1994;
06-JUN-1995;
     New monoclonal antibody to human interleukin-5 - used to produce products for the treatment and diagnosis of conditions associated with excess eosinophil prodn., e.g asthma etc.
                                     N-PSDB; AAT34094
                                                                                   (SMIK )
                                                                                                                                                          11-JUL-1996
                                                                                                                                                                                                                                                                                                                                               interleukin-5; II-5; eosinopnii; asummu, ____atopic dermatitis; therapy; diagnosis; light chain; VL; monoclonal antibody; MAb.
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                                                                                                                                         22-DEC-1995;
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                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                      Antibody engineering; humanised antibody; chimeric antibody; interleukin-5; IL-5; eosinophil; asthma; allergic rhinitis;
                                                                                                                                                                                                                                                                                                                                                                                                 Humanised
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                                             1996-333976/33
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                                                                                  SMITHKLINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18; Conser
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                                                            Appelbaum ER, McMillan LJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                          95US-0470110.
94US-0363131.
95US-0467420.
                                                                                                                                                                                                                                                                                                                                                                                              antibody light chain variable region.
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95..10:
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63..94
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56..62
                                                                                                                                                                                                                                                                            /note=
41..55
                                                                                                                                                                                               104..113
                                                                                                                                                                                                      /note= "mouse MAb
                                                                                                                                                                                                                                                                                                           /note-
                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                               /label= CDR3
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                                                                                   BEECHAM O
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100.0%
                                                                                                                                                                                                                            "human
                                                                                                                                                                                                                                            "mouse
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                                                                                                                                                                                                                                                                   "human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       against Plasmodium infections.
                                                           Chaiken IM,
Theisen TW;
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Pred. No.
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                                                                                                                                                                                       framework region"
                                                                                                                                                                                                                                                                    framework region"
                                                                                                                                                                                                                                                                                   286
                                                                                                                                                                                                      2B6 VL CDR3"
                                                                                                                                                                                                                             framework region"
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                                                                                                                                                                                                                                                                                   VL CDR1"
                                                                    Cook RM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 15,
5. 7.8e-05;
0;
                                                                    Gross MS;
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ID AAW4
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Best Local S
Matches 18
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                                                                                                               24 - DEC - 1997
                                                                                                                                                          WO9748418-A1
                                                                                                                                                                                                                                               Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Neutralising antibody; monoclonal antibody; MAb; 2B6; LEN; mouse; interleukin-5; IL-5; human; eosinophil; asthma; allergic rhinitis; atopic dermatitis; therapy; diagnosis; humanised antibody.
                                                                 20-JUN-1997;
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18; Conserv
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/note- "2B6
104..113
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63..94
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100.0%; Pr
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Pred. No.
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7.8e-05;
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Best Local Similarity
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Polynucleotides encoding anti-p53 antibodies, fragments, useful in treatment and diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Appelbaum ER,
                                                                                        N-PSDB; AAA96134.
                                                                                                                                                                                                                                                                                                           19-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                p53; antibody; immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Amino acid sequence of anti-p53 antibody light chain clone 163.2.
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                                                                                                                      WPI; 2000-638249/61
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                                                                                                                                                                                    Coomber DWJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               response; vaccine; gene therapy; coronary heart disease.
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Pred. No.
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   of cancer, rheuman
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC an immune response against a naturally occurring disease. The antibodies CC are useful in pharmaceutical compositions, which additionally contain CC chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes CC or gadolinium. The polypeptide components of the antibodies are useful in CC varcines, for inducing an immune response against a disease in a CC vertebrate, for treatment and/or prophylaxis of disease and for detection CC purposes. The nucleic acid sequences can be used to detect a disease as CC well as for gene therapy and recombinant production of the polypeptides. CC in particular, the following can be treated cancer, rheumatoid arthritis CC and coronary heart disease. Cancers include carcinogenic tumours, tumours CC of epithelial origin, e.g. colo-rectal cancer, breast cancer ung cancer, CC head and neck tumours, hepatic cancer, prostate cancer and CC gastric cancer, brain cancer, bladder cancer, mesenchymal tumours, CC e.g. sarcoma, and hemopoletic tumours, e.g. B cell lymphoma.
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Best Local Similarity
                                                                                     Polynucleotides encoding anti-p53 antibodies, fragments, useful in treatment and diagnosis carthritis and coronary heart disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents the light chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 30;
                                                                                                                                                     N-PSDB;
                                                                                                                                                                                                    Ward RL,
                                                                                                                                                                                                                                                               19-MAR-1999;
                                                                                                                                                                                                                                                                                             15-MAR-2000; 2000WO-AU00189
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                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                    p53; antibody; immune response; vaccine; gene therapy; cancer;
rheumatoid arthritis; coronary heart disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amino acid sequence of anti-p53 antibody light chain clone 163.6.
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hes 0;
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                                                                                                         of cancer, rheumatoid
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The present sequence represents the light chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies

Claim

30; Page 136; 163pp; English.

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                      The present sequence represents the light chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes
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                                                                                                                                                                                                                Claim
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hritis; coronary heart disease.
ys, prodrugs, toxins and imaging markers e.g. radioisotor The polypeptide components of the antibodies are useful
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Pred. No.
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7.8e-05;
                                                                                                                                                                                                                                                                                                of cancer, rheumatoid
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Best Local
                 an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in vaccines, for inducing an immune response against a disease in a vertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as well as for gene therapy and recombinant production of the polypeptides. In particular, the following can be treated cancer, rheumatoid arthritis and coronary heart disease. Cancers include carcinogenic tumours, tumours
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Pred. No.
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7.8e-05;
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                                   The present sequence represents the light chain of an antibody reactive CC against p53. The antibody is obtained from a vertebrate host expressing CC an immune response against a naturally occurring disease. The antibodies CC are useful in pharmaccuttoal compositions, which additionally contain CC chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes CC or gadolinium. The polypeptide components of the antibodies are useful in CC vertebrate, for inducing an immune response against a disease in a CC vertebrate, for treatment and/or prophylaxis of disease and for detection CC purposes. The nucleic acid sequences can be used to detect a disease as CC well as for gene therapy and recombinant production of the polypeptides. CC in particular, the following can be treated cancer, rheumatold arthritis CC and coronary heart disease. Cancers include carcinogenic tumours, tumours CC pastric cancer, brash cancer, bapatic cancer, presst cancer lung cancer, CC thead and neck tumours, hapatic cancer, panorealic cancer, ovarian cancer, brashric cancer, brash cancer, bandder cancer, mesenchymal tumours, CC essettic cancer, brash cancer, brash cancer, cancer lung cancer, cancer lung cancer, cancer lung cancer, cancer, brash cancer, brash cancer, cancer lung cancer, cancer, cancer lung cancer, cancer, brash cancer, cancer, brash cancer, cancer, cancer lung cancer, can
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rheumatoid arthritis; coronary heart disease.
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DB 21; . 7.8e-05;

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                      The present sequence represents the light chain of an antibody reactive cagainst p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies care useful in pharmaceutical compositions, which additionally contain challenges, from the products, toxins and imaging markers e.g. radioisotopes cor gadolinium. The polypeptide components of the antibodies are useful in contract, for inducing an immune response against a disease in a current card and for prophylaxis of disease and for detection contracts. For uncleid acid sequences can be used to detect a disease as compared to the polypeptides. In particular, the following can be treated cancer, rheamatold arthritis and coronary heart disease. Cancers include carcinogenic tumours, tumours of epithelial origin, e.g. colo-rectal cancer, breast cancer lung cancer, cancer badder cancer, postate cancer, ovarian cancer, based nand neck tumours, hepatic cancer, prostate cancer, ovarian cancer, cancer bladder cancer, mesenchymal tumours, ce.g. sarcoma, and hemopoietic tumours, e.g. B cell lymphoma.
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Best Local S
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Sequence
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Db 71 sgsgtdftltiss1qaed 88

Search completed: July 15, 2002, 12:57:59
Job time: 415 sec

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Title:
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Maximum Match 10
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    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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   protein search, using sw model
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Listing first 45 summaries
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(c) 1993 - 2000 Com
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US-08-667-769A-21
US-08-667-769A-21
US-08-637-647-21
US-08-525-539A-86
US-09-014-880-6
US-09-017-628-6
US-09-017-628-6
US-09-025-769B-17
US-09-025-539A-65
US-08-483-636-14
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Query Matc Best Local Matches	RESULT 1 US-08-554-840-1 Sequence 13, Patent No. 60 GENERAL INFO APPLICANT: CORRESPOND ADDRESSE STATE: COUNTRY: ZIP: 22 COMPUTER R MEDIUM T COMPUTER R OPERATING SOFTWARE CURRENT APPLICAT FILING D CLASSIFI ATTORNEY/A NAME: T REGISTRA REFERENC TELEFOMMUN TELEFAX: INFORMATION SEQUENCE C LENGTH: TYPE: a STRANDED TOPOLOGY NOLEGULE T US-08-554-840-1	444440087 5443008
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60.4%; 100.0%; ive (US/08554 bil bil duardo A. cland A. HUMANIZED COMPOSITI S: Doane, Sw 404 404 404 404 404 404 404 404 404 40	133 3 US- 133 4 US- 135 3 US- 155 3 US- 171 4 US- 260 4 US- 275 4 US- 275 4 US- 285 3 US- 342 4 US- 495 4 US- 495 4 US-
Score 87; DB 3; Pred. No. 1.1e-); Mismatches	ANTIBODIES TO ONS CONTAINING ecker & Mathis +1.0, Version + 54,840	-08-463-903-4 -07-935-695-4 -08-812-586-46 -08-828-7418-11 -09-160-567-11 -09-160-567-11 -09-16-98A-56 -07-935-695-20 -07-935-695-2 -07-935-695-2 -07-935-695-6 -08-463-903-6 -08-463-903-6 -08-463-903-6 -08-463-903-6 -08-463-903-6 -08-463-903-6 -08-463-903-6 -08-463-903-6 -08-463-903-6 -08-463-903-6 -08-463-903-6 -09-160-567-6 -09-160-567-6 -09-160-567-4 -09-160-567-4
Length 80; 05; 0; Indels	HUMAN 9P39, AND THERAPEUTIC	Sequence
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Gaps (THEREOF	4, Appli 4, Appli 11, Appli 11, Appl 20, Appl 50, Appli 2, Appli 6, Appli 6, Appli 6, Appli 6, Appli 6, Appli 6, Appli 7, Appli 6, Appli 7, Appli 7, Appli 8, Appli 9, Appli
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; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-554-840-15
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FILING DATE: 07-NOV-1995
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35,030
REFERENCE/DOCKET NUMBER: 012712-127
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-6620
                                                                                                                                GENERAL INFORMATION:
APPLICANT: BLACK, Amelia
APPLICANT: HANNA, Nabil
APPLICANT: PADLAN, Eduaro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                   APPLICANT: NEWMAN, ROLAND A. TITLE OF INVENTION: HUMANIZED ANTIBODIES TO HUMAN 9p39, TITLE OF INVENTION: COMPOSITIONS CONTAINING AND THERAPP NUMBER OF SEQUENCES: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                   CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION: COMPOSITIONS
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APPLICANT: HANNA, Nabil
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CITY: Alexandria
STATE: Virginia
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les 18; Conserv
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                ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SGSGTDFTLTISSLQAED 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        i: 80 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22313-1404
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                                                                                                                              PADLAN, Eduardo A.
                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
Burns, Doane, Swecker & Mathis
O. Box 1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        single
                                                                                                                                                                                                                                                                                                                                                                                                   60.4%; Score 87; DB 3; 100.0%; Pred. No. 1.1e-(tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28
                                                                                                                                                                                                                                                                                                                                                                                                     1.1e-05;
hes 0;
                                                                      AND THERAPEUTIC USE THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 80;
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US-08-525-539A-81
; Sequence 81, Application US/08525539A
; Patent No. 6309636
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                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: I
APPLICANT: I
APPLICANT: I
ATTORNEY/AGENT INFORMATION:
NAME: DYLAN, TYLER
REGISTRATION NUMBER: 37,612
REFERENCE/DOCKET NUMBER: 27633-20001.21
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-6620
                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 81
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Teskin, Robin L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PATENTIN Rela
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 07-NOV-199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                       STREET:
                                                                                     FILING DATE: 14-SEP-1995
CLASSIFICATION: 424
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                                                                                                                                                                                                                                                                   COUNTRY:
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                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                    ZIP: 94304-1018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            65 SGSGTDFTLTISSLQAED 82
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: United States ZIP: 22313-1404
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STATE: Virginia
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                                                                                                                                                                                                                                                                                  Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                     USA
                                                                                                                                                                                                                                                                                                                                                                                                                                              PETERSON, JERRY A.
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100.0%;
                                                                                                                                                                                                                                                                                                                                                                                           RECOMBINANT PEPTIDES DERIVED FROM THE MC3 ANTI-BA46 ANTIBODY, METHODS OF USE THEREOF, METHODS OF HUMANIZING ANTIBODY PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Release #1.0, Version #1.30
                                                                                                                           US/08/525,539A
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Pred. No.
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                                                                                                                                                            Version
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1.5e-05;
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; MOLECULE TYPE: peptide
US-07-942-245-30
RESULT 6
US-08-467-420A-21
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                                                                                                                                                                                                                                          APPLICATION NUMBER: US/07/942.

APPLICATION NUMBER: US/07/942.

FILING DATE: 09-SEP-1992

CLASSIPICATION: 530

TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 293-7060
TELEFAX: (202) 293-7860
TELEFX: 6491103
INFORMATION FOR SEQ ID NO: 30:
SEQUENCE CHARACTERISTICS:
LENGTH: 112 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 30
Patent No.
                                                                                                      Query Match
Best Local Similarity 100.
Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: ROGUSKA,
APPLICANT: GUILL,
TITLE OF INVENTION: S
TITLE OF INVENTION: 2
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: HP 9000/700 Workstation
OPERATING SYSTEM: UNIX
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LENGTH: 107 amino acids
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CITY: Washington
STATE: D.C.
                                                                         12 SGSGTDFTLTISSLQAED 29
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STRANDEDNESS: sin
TOPOLOGY: linear
                                                           71
                                                                                                                                                                                                                                    TYPE: amino acid
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                                                         SGSGTDFTLTISSLQAED 88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TRY: United States 20037-3202
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706141
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEARLE,
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100.0%; Pred. No. 1.5e-C
vative 0; Mismatches
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Anthony R.
Anthony R.
Michael A.
Braydon C.
SURFACE RESIDUE VENEERING OF RODENT
ANTIBODIES
522
                                                                                                                  60.4%; Score 87; DB 1; I
100.0%; Pred. No. 1.5e-05;
tive 0; Mismatches 0;
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Sequence 21, Application US/08470110A
Patent No. 5693323
GENERAL INFORMATION:
APPLICANT: Ames, Robert S.
APPLICANT: Appelbaum, Edward R.
APPLICANT: Chaiken, Irwin M.
APPLICANT: Cook, Richard M.
APPLICANT: Gross, Mitchell S.
APPLICANT: Holmes, Stephen D.
APPLICANT: Holmes, Stephen D.
                                                                                                                                                                                        US-08-470-110A-21
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                                                                                                                                                                                                               RESULT
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Ames, Novelland, Edward APPLICANT: Appelbaum, Edward NobilCANT: Chalken, Irwin M.
                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 610 270-5024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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ADDRESSEE: SmithKline Beecham Corp./Corporate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                71 SGSGTDFTLTISSLQAED 88
                                                                                                                                                                                                                                                                                                   12 SGSGTDFTLTISSLQAED 29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 19406-0939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: P. O. Box 1539-UW2220
CITY: King of Prussia
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                       Local Similarity
les 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Intellectual Property
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            113 amino acids
                          Cook, Richard M.
Gross, Mitchell S.
Holmes, Stephen D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                McMillan, Lynette J
Theisen, Timothy W.
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Holmes, Stephen D.
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McMillan, Lynette J
Theisen, Timothy W.
                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                  60.4%; Score 87; DB 1; Le 100.0%; Pred. No. 1.6e-05; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Treatment of IL5 Mediated Disorders
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                                                                                                                                                                                                                                                                                                                                                                    Length 113;
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TITLE OF INVENTION: RETITLE OF INVENTION: Tr NUMBER OF SEQUENCES: 7 CORRESPONDENCE ADDRESS:

Recombinant IL5 Antagonists Useful in Treatment of IL5 Mediated Disorders

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APPLICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
ATTORNEY_AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: SUTTON, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERRACE/DOCKET NUMBER: P50
TELECOMMUNICATION INFORMATION:
TELECHONE: 610 270-5024
                                                                                             NUMBER OF SEQUENCES: 76
CORRESPONDENCE ADDRESS:
ADDRESSEE: Smithkline Beecham Corp./Corporate
                                                                                                                                TITLE OF INVENTION: Recombinant IL5 Antagonists Useful in TITLE OF INVENTION: Treatment of IL5 Mediated Disorders NUMBER OF SEQUENCES: 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: protein
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COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: single
                                COUNTRY:
                                                STATE:
                                                              CITY: King of Prussia
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nes 18; Conserv
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FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                     21, Application US/08667769A
o. 5783184
                   19406-0939
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Gross, Mitchell S.
Holmes, Stephen D.
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                                  USA
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                                                                                  Box 1539-UW2220
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100.0%; pr
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; Pred. No. 1.6
0; Mismatches
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US-08-940-371-21
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 610-270-5090 INFORMATION FOR SEQ ID NO:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                               APPLICANT:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5024
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: PCT/
FILING DATE: 22-DEC-1995
                                                                                                                                                         NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                       APPLICANT: McMillan, Lynette J.
APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinant IL5 Antagonists Useful in
TITLE OF INVENTION: Treatment of IL5 Mediated Disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: protein
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APPLICATION NUMBER: US 08/467420
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APPLICATION NUMBER:
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CLASSIFICATION: 424
                                                                                    STREET: P. O. Box 1539-UW2220
CITY: King of Prussia
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                                     COUNTRY: USA
ZIP: 19406-0939
                                                                                                                       ADDRESSEE: SmithKline Beecham Corp./Corporate ADDRESSEE: Intellectual Property
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE:
                                                                         STATE:
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Holmes, Stephen D.
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06-JUN-1995
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100.0%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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PILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
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                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM CC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                               ADDRESSEE: Intellectual Property STREET: P. O. Box 1539-UW2220 CITY: King of Prussia
                                                                  CLASSIFICATION:
                                                                            APPLICATION NUMBER: US/08/637,647
FILING DATE: 19-JUN-1997
                                                                                                                                                                                                                COUNTRY: USA
ZIP: 19406-0939
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TOPOLOGY: 111
CLASSIFICATION:
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les 18; Conserv
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SOFTWARE: PatentIn Release #1.0, Version #1.25
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Recombinant IL5 Antagonists Useful in Treatment of IL5 Mediated Disorders
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                                US/08/363,131
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; Pred. No. 1.6e-05;
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US-08-525-539A-80
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acid
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TOPOLOGY: line
MOLECULE TYPE: |
                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/01
FILLING DATE: 14-SEP-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: DYLAN, TYLER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDLIM TYPE: Floppy disk
COMPUTER: IBM PC COMPUTER: IBM PC COMPUTER: IBM PC COMPUTER. IBM PC COMPUTER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 610 270-5024
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NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
                                                                                                                                                                    SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
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TITLE OF INVENTION:
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                                                                                                                                                                                                                                           TELEFAX: (415
TELEX: 706141
                                 STRANDEDNESS:
TOPOLOGY: li
                                                                                                    TYPE:
                                                                                                                                                                                                                                                                                                       TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 37,612
REFERENCE/DOCKET NUMBER: 27
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STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12 SGSGTDFTLTISSLQAED 29
                                                                                                                                  LENGTH:
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(415) 494-0792
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100.0%; Pr
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Pred. No.
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Query Match Best Local Similarity

60.4%; 100.0%;

Score 87; Pred. No.

DB 4; I

Length 113;

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                                                                                                                 RESULT 13
PCT-US93-08435-8
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                                                                                   Sequence 8, Application PC/TUS9308435 GENERAL INFORMATION:
                                                                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (215) 540-9200
TELEFAX: (215) 540-5818
INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DCS/MS-DOS
SOTTWARE: PATENTIN RC-DEASE #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/08435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0'
FILING DATE: 09-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS
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REFERENCE/DOCKET NUMBER: SBC P50107
TELECOMMUNICATION INFORMATION:
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                              APPLICANT:
                                            AFPLICANT:
                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein
               APPLICANT:
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CITY: Spring House
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                                                                                                                                                                            SGSGTDFTLTISSLQAED 88
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OF INVENTION: Novel Antibodies for Conferring Passive
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                  the Navy
U. S. Government, Secretary of
                                                        U. S. Government,
              the Army
                                                                        SmithKline Beecham, Corporation
                                                                                                                                                                                                                                     Conservative
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Pred. No.
                                                          Secretary of
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1.6e-05;
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RESULT 14
PCT-US95-17082A-21
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APPLICANT:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
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REFERENCE/DOCKET NUMBER: SB
TELECOMMUNICATION INFORMATION:
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APPLICATION NUMBER: PCT/US93/08435
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                                                                                                                                                                                NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                   TITLE OF INVENTION: Treatment of IL5 Mediated Disorders
                                                                                                                                                                                                                             APPLICANT: Theisen, Timothy W. TITLE OF INVENTION: Recombinan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12 SGSGTDFTLTISSLQAED 29
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hes 18; Conservat
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CLASSIFICATION:
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                                                                                                                                ADDRESSEE: SmithKline Beecham Corp./Corporate STREET: P.O. Box 1539-UW2220 CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      71 SGSGTDFTLTISSLQAED 88
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Box 457, 321 Norristown Road
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                                                                                                                                                                                                                                                                                                    Gross, Mitchell S.
                                                                                                                                                                                                                                                                                                                                  Appelbaum, Edward
Chaiken, Irwin M.
                                                                                                                                                                                                                                                                                                                                                                       Ames,
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                                                                                                                                                                                                                                                                 McMillan, Lynette J.
                                                                                                                                                                                                                                                                                     Holmes, Stephen D.
                                                                                                                                                                                                                                                                                                                  Cook, Richard M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                       Robert S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       540-5818
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                                                                                                                                                                                                                                      Recombinant IL5 Antagonists Useful in
 Release #1.0,
                                                                                                                                                                                                                                                                                                                                                    Edward R
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     Version #1.30
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/17082A

FILING DATE

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; TYPE: amino acid
; STRANDEDNESS: Single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
PCT-US95-17082A-21
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FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: SUTTON, JEffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P502
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 21:
        FILING DATE:
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/905
FILING DATE: June 29, 1992
                                                                                                                                                                                                                                                                                                                                        APPLICANT: No. 5767246ihiko ITO
APPLICANT: Kazuhiro NAGAIKE
TITLE OF INVENTION: Human Monocl
TITLE OF INVENTION: Specifically
TITLE OF INVENTION: Cell Membran
                                                                                                            SOFTWARE: WordPerfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.
COMPUTER: IBM Compatible
OPERATING SYSTEM: MS-DOS
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ADDRESSEE: Wenderoth,
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                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
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                                                                                              APPLICATION NUMBER:
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APPLICATION NUMBER:
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                              07/905,534
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Best Local Similarity
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OTHER INFORMATION:
PUBLICATION INFORMATION:
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CHROMOSOME/SEGMENT:
MAP POSITION:
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ORIGINAL SOURCE:
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REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-8850
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IMMEDIATE SOURCE:
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HYPOTHETICAL:
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FILING DATE:
PUBLICATION DATE:
RELEVANT RESIDUES
                                                                          12 SGSGTDFTLTISSLQAED 29
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VOLUME:
                                                                                                                                                                                                                                                                                                                                                               NAME/KEY:
LOCATION:
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CELL TYPE:
CELL LINE:
                                                          71 SGSGTDFTLTISSLQAED
                                                                                                                                                                                                                                                                                                  TITLE:
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TISSUE TYPE:
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Pred. No. 1.6e-05;
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Perfect score:
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        protein search, using sw model
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2: pir2:*
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        July 15, 2002, 13:01:03; Search time 95.45 Seconds (without alignments) 29.194 Million cell updates/sec
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107	107	107	105	104	104	104	104	104	104	103	103	103	103	103	102
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D30535	G30535	B3053	A3053	C3053	PH1104	PH1103	PH1102	PH110:	PH1053	PH1047	PH1054	PH1052	PH1051	PH1050	A34153
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ALIGNMENTS

RESULT S37533

Ig kappa chain V region (V-kappa 4) - human (fragment)
C.Species: Homo sapiens (man)
C.Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_
C.Accession: S37533
R.Klein, U.; Kueppers, R.; Rajewsky, K.
submitted to the EMBL Data Library, September 1993
A.Description: Human IgM(+)IgD(+) cells, the major B cell
A; Reference number: S37501
A; Reference number: S37501
A; Accession: S37533
A; Status: preliminary
A; Molecule type: mRNA
A; Residues: 1-92 cKLE>
A; Crearing and control of the control of

cell subset

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06-Jan-1995 #text_change

23-Jul-1999

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Ig kappa chain V region (V-kappa 4) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
C;Accession: S37530
R;Klein, U; Kueppers, R: Rajewsky, K.
submitted to the EMBL Data Library, September 1993
A;Description: Human 1gM(+)IgD(+) cells, the major B cell subset in the periphe
A;Reference number: S37501
A;Accession: S37501
A;Accession: S37501
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-92 <KLE>
A;Residues: 1-92 <KLE>
A;Residues: 1-92 <KLE>
A;Cross-references: EMBL: Z36630; NID:9405702; PIDN:CAA81383.1; PID:9405703
C;Superfamily: Immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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                                 Score 63; DB Pred. No. 0.31 0; Mismatches
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Ig kappa chain V region (V-kappa 4) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
C;Accession: S37531
R:Klein, U:; Kueppers, R.; Rajewsky, K.
submitted to the EMBL Data Library, September 1993
A;Description: Human IgM(+)IgD(+) cells, the major B cell subset in the per A;Reference number: S37501
A;Accession: S37531
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ig kappa chain V region (V-kappa 4) - human (fragment)
C;Specics: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
C;Accession: S37535
R;Klein, U; Kueppers, R.; Rajewsky, K.
submitted to the EMBL Data Library, September 1993
A;Description: Human IgM(+)IgD(+) cells, the major B cell subset in the per:
A;Reference number: S37501
A;Accession: S37535
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submitted to the EMBL Data Library, September 1993
A;Description: Human IgM(+)IgD(+) cells, the major B cell subset in the peripheral blood A;Reference number: S37501
A;Accession: S37529
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-92 <KLE>
A;Cross-references: EMBL:Z26631; NID:g405700; PIDN:CAA81384.1; PID:g405701
C;Superfamily: Immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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C:Species: Homo sapiens (man)
C:Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
C:Accession: S37529
R:Klein, U.: Kueppers, R.: Rajewsky, K.
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S37531
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A:Molecule type: mRNA
A:Molecule type: mRNA
A:Residues: 1-92 <KLE>
A:Cross-references: EMBL:Z26625; NID:g405712; PIDN:CAA81378.1; PID:g405713
C:Superfamily: immunoglobulin V region; immunoglobulin homology
C:Keywords: heterotetramer; immunoglobulin
A; Status: preliminary A; Molecule type: mRNA
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Pred. No. 0.31;
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A;Cross-references: EMBL:226628; NID:9405706; PIDN:CAA81381.1; C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin
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C;Accession: S37532
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C:Species: Homo sapiens (man)
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A;Residues: 1-92 KLED.
A;Residues: 1-92 KLED.
A;Cross-references: EMMI: 226626; NID: g405710; PIDN: CAA81379.1; PID: g405711
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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C;Accession: S37534

C;Accession: S37534

R;Klein, U.; Kueppers, R.; Rajewsky, K.

Submitted to the EMBL Data Library, September 1993

Submitted to the EMBL Data Library, September 1993

A;Description: Human IgM(+)IgD(+) cells, the major B cell subset in A;Reference number: S37501

A;Accession: S37534
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37534
Says chain V region (V-kappa 4) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 23-Jul-1999
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A; Residues: 1-92 <KLE>
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RESULT 10
A49138
IgA kappa rheumatoid factor variable - human C; Specles: Homo sapicns (man)
C; Date: 19-Dec-1993 #sequence_revision 18-No C; Accession: A49138
R; Gause, A.; Kuppers, R.; Mierau, R. Clin. Exp. Immunol. 88, 430-434, 1992
A; Title: A somatically mutated V kappa IV ge A; Accession: A49138
A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ig kappa chain V-IV region (Fue) - human C;Species: Homo sapiens (man) C;Date: 11-Apr-1990 *sequence_revision 11-Apr-1990 *text_change 21-Jan-2000 C;Accession: B34153 R;Mihaesco, E.; Ayadi, H.; Congy, N.; Gendron, M.C.; Roy, J.P.; Heyermann, H.; Frangio J. Biol. Chem. 264, 21481-21485, 1989 A;Title: Multiple mutations in the Wariable region of the kappa light chains of three A;Reference number: A34153; MUID:90094313 A;Accession: B34153
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A; Molecule type: DNA
A; Mesidues: 1-101 <MAN>
A; Note: residues 28-33 were obtained from Figure 4
C; Comment: This antibody is produced by Epstein-Barr virus-transformed B cell that bears C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 1-23/Region: framework 1
F; 16-56/Domain: immunoglobulin homology < IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Molecule type: protein
A;Residues: 1-102 <MITHO
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;16-96/Domain: immunoglobulin homology <IMM>
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F;63;94/Region: framework 3
F;95-101/Region: complementarity-determining
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A;Title: Molecular characteristics of antibodies bearing an anti-DNA-associated idiotype A;Reference number: PH0862; MUID:92078875
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C:Species: Homo sapiens (man)
C:Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 21-Jan-2000
C:Accession: PH0869
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F; 41-55/Region:
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complementarity-determining 1
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                                                                                                                                                   18-Nov-1994 #text_change 21-Jan-2000
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A:Molecule type: DNA
A:Residues: 1-106 <GAL>
A;Residues: 1-106 <GAL>
A;Residues: 1-106 <GAL>
A;Residues: PID:g298207; PIDN:AAB22366.1; PID:g298208
A;Note: sequence inconsistent with the nucleotide translation
A;Note: sequence extracted from NCBI backbone (NCBIN:106633, NCBIP:106637
C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;20-99/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ig kappa chain V region - hum
C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence
C;Accession: S30523
                                                                                                                                  A;Cross-references: EMBL;218325
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Reywords: heterotetramer; immunoglobulin
F;16-96/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                       Ig kappa chain V region - human C;Species: Homo sapiens (man) C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change C;Dates: 05-Jan-1995 #sequence_revision 06-Jan-1995 #text_change C;Accession: S30520
QY
                                                                                                                                                                                                                      A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-113 <MAR>
                                                                                                                                                                                                                                                                                A; Accession: S30520
                                                                                                                                                                                                                                                                                                 submitted to the EMBL Data Library, A; Reference number: $30520
                                                                                                                                                                                                                                                                                                                                          R; Mariette, X.
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
E;16-96/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: mRNA
A; Residues: 1-113 <MAR>
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A; Reference number: S30520
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                                      Score 63; DB 2;
Pred. No. 0.37;
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SGSGTDFTLTISSLQAED

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Ig kappa chain V-IV region (Len) - human
C:Species: Homo saplens (man)
C:Cacesion: AO1903; F61458
C:Accession: AO1903; F61458
R;Schneider, M.; Hilschmann, N.
Hoppe-Seyler's Z. Physiol. Chem. 356, 507-557, 1975
A:Title: Die Primaerstruktur einer monoklonalen Immunglobulin-L-Kette der Subgruppe IV:A;Reference number: AO1903; MUID:76004342
A;Reference number: AO1903; MUID:76004342
A;Accession: AO1903
A;Molecule type: protein
A;Residuks: 1-114 <SCH>
A;Note: this is the first completely sequenced V region of a new kappa chain subgroup, A;Note: this is the first completely sequenced V region of a new kappa chain subgroup, A;Note: the C region of this chain has the Inv (3) marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ig kappa chain V region - human
C;Species: Homo sapiens (man)
C;Date: 02-Dec:1993 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
C;Accession: S34003
R;Mariette, X; Tsapis, A.; Brouet, J.C.
Eur. J. Immunol. 23, 846-851, 1993
A:Title: Nuclectidic sequence analysis of the variable domains of four human A;Reference number: S34001; MUID:93209281
A;Accession: S34003
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-113 <MARP
A;Residues: 1-113 <MARP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: EMBL:Z18328
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;16-96/Domain: immunoglobulin homology <IMM>
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Pred. No. 0.37;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Brouet, J.C.; Dellagi, K.; Gendron, M.C.; Chevalier, A.; Schmitt, C.; Mihaesco, E. J. Exp. Med. 170, 1551-1558, 1989
A;Tille: Expression of a public idiotype by human monocional IgM directed to myelin-a A;Reference number: A61458; MUID:90039128
  맑
                                                                                                                                                                                                                                                                                                                                               C; Comment: This is a Bence Jones protein.
                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: protein
A; Residues: 1-13 <BRO>
                                                                                                                                                                                                                                                                                                                                                                                                                      A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Accession: F61458
                                                                                           Matches
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Best Local Similarity
  71
                      12 SGSGTDATAPASSLQAED 29
SGSGTDFTLTISSLQAED
                                                                                        14;
                                                                                           Conservative
                                                                                                               43.8%;
77.8%;
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Pred. No. 0.37;
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Search completed: July 15, 2002, 13:01:03 Job time: 469 sec

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Result
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(c) 1993 - 2000 Comp
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KV4B_HUMAN
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             KYJE_HUMAN
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TAT_HVIBR
TAT_HVIBR
TAT_HVIZ
TAT_HVIZ
TAT_HVIZ
TAT_HUZ
TAT_HUZ
TAT_HUZ
KYJA_HUMAN
KYJA_HUMAN
RPC3_YEAST
TAT_HVIAZ
TAT_HVIAZ
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KV3J_HUMAN
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RESULT 2
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Qy

12 SGSGTDATAPASSLOAED SGSGTDFTLTISSLQAED

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Query Match Best Local S Matches 14

14; Conservative

43.8%;

Score 63; DB Pred. No. 0.08 0; Mismatches

DB 1;

Length 114; Indels

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Gaps

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COMPLEMENTARITY-DETERMINING-2. FRAMEWORK-3. COMPLEMENTARITY-DETERMINING-3. FRAMEWORK-4. BY SIMILARITY.

COMPLEMENTARITY-DETERMINING-1. FRAMEWORK-2.

DISULFID NON_TER SEQUENCE DOMAIN

114 AA;

114 12640 MW;

0647F1D17F236485 CRC64;

Immunoglobulin 1
DOMAIN 1
DOMAIN 24
DOMAIN 41
DOMAIN 56
DOMAIN 63
DOMAIN 95

ion;

Bence-Jones protein. FRAMEWORK-1.

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                                                                                                                                                                                                                                      MEDLINE-86041854; PubMed-2997713; Marsh P., Mills F., Gould H.; "Detection of a unique human V ka cDNA probe.";
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Blochem. J. 171:337-347(1978).

Blochem. J. 171:337-347(1978).

HISCELLANEOUS: THE MATURE CHAIN HAS 12 ADDITIONAL RESIDUES AT TTS AMINO END, DUE TO A TANDEM DUPLICATION OF 36 NUCLEOTIDES AFTER THE CODON FOR RESIDUE 36. RESIDUE 42 CORRESPONDS TO THE AMINO-TERMINAL CODON FOR RESIDUE 36. RESIDUE 42 CORRESPONDS TO THE AMINO-TERMINAL
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MEDLINE*78186617; PubMed*418775;
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                                                                                                                                                                                                                                                                                                                                                             SEQUENCE OF 41-149 FROM N.A. MEDLINE-80176554; PubMed-6245773;
                                                                                                                                                                                                                                                                                                                                                                                                                             5'-terminal regions of immunoglobulin variable-region Cell 29:681-689(1982).
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kelley D.E., Coleclough C., Perry R.P.; Functional Significance and evolutionary development 5'-terminal regions of immunoglobulin variable-region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE OF 1-71 FROM N.A. MEDLINE=83001944; Pubmed=6288267;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse).
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SMART; SM00406; IGV;
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PIR; A01905; K4HU17.
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                                                                                                                                                                                             Smith G.P.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ig kappa chain V-V region MPC11 precursor.
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                                                                                                                                                                  "Sequence of the full-length immunoglobulin kappa-chain
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an email to license@isb-sib.ch).
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(Rel. 01, Last sequence update)
(Rel. 38, Last annotation update)
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COMPLEMENTARITY-DETERMINING-2
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Pfam; PF00047; 1g; 1.

SMART; SM00406; IGv; 1.

Immunoglobulin V region; Bence-Jones protein.

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                                                                                                                                InterPro; IPR003596;
                                                                                                                                                          PIR; A01868; K1HUHU.
HSSP; P80362; 1WTL.
                                                                                                                                                                                                                     subgroups.
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15-JUL-1999
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                                           FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
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Pred. No. 0.38;
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COMPLEMENTARITY-DETERMINING-3
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COMPLEMENTARITY-DETERMINING-2.
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                                                                                                                                                                                                                                                                                                                 DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entities requires a license agreement (S or send an email to license@isb-sib.ch).
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13-AUG-1987 (Rel. 05, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
19 kappa chain V-III region VH precursor (Fragment).
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SMART; SM00406; IGV;
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c Acids Res. 12:9229-9236(1984).
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88 | 1
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11671 MW;
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72.2%;
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BY SIMILARIT
                                                                                                      Score 58; I
Pred. No. 0
                                                                                                                                                                                                                                                                                                              IG KAPPA CHAIN V-III REGION VF
FRAMEWORK-1.
COMPLEMENTARITY-DETERMINING-1.
FRAMEWORK-2.
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Pred. No. 0.36;
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                                                                                                                                                                                        51CD55BA53B21929 CRC64
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                                                                                                                                                                                                                               SIMILARITY
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KYII_HUMAN
ID KVII_HUMAN
AC PO1601;
DT 21_JUL-1986

RESULT 9
KV1W_HUMAN
ID KV1W_H
AC P04431
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Best Local
    KV1W_HUMAN P04431;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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BentLey D.L., Rabbitts T.H.;
"Human immunoglobulin variable region
kappa genes_and_a_pseudogene.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; ig; 1.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the El
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  conversion.";
Cell 32:181-189(1983).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bentley D.L., Rabbitts T.H.; "Evolution of immunoglobulin V genes: evidence indicating recently duplicated human V kappa sequences have diverged
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A. MEDLINE-83129397; PubMed-6402305;
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HSSP; P01607; 1REI.
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                        STANDARD;
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01, Last sequence update)
38, Last annotation update)
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Pred. No.
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ERAMEWORK-1.
ERAMEWORK-2.
ERAMEWORK-2.
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                        PRT;
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by gene
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RESULT 10
TAT_HV112
ID TAT_HV
AC P04326
AC P04326
DT 10-OAF
DT 13-AUG
DT 13-AUG
DT TAT_F
CON TAT
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Best Local
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P04326;
20-MAR-1987
13-AUG-1987
16-OCT-2001
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-AUG-1987
13-AUG-1987
15-JUL-1999
Arya S.K., Gallo R.C.; "Three novel genes of human T-lymphotropic virus type III: reactivity of their products with sera from acquired immune
                                                                    SEQUENCE FROM N.A. MEDLINE=86177573; PubMed=3008154;
                                                                                                                                  Viruses; Retroid
NCBI_TaxID=11679;
                                                                                                                                                                                                                     TAT protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00047; ig; 1. SMART; SM00406; IGV;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A. MEDLINE-85014148; PubMed-6091049;
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                                                                                                                                                                            Human immunodeficiency virus type 1
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(Rel. 05, Last sequence update)
(Rel. 40, Last annotation update)
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05, Last sequence update)
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                                                                                                                                                       Retroviridae;
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Pred. No.
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BY SIMILARITY.
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FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2
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FRAMEWORK-1
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                                                                                                                                                                                                                                                                                                                                    PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               KAPPA CHAIN V-I REGION WALKER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              precursor.
                                                                                                                                                                               (clone 12) (HIV-1).
                                                                                                                                                                                                                                                                                                                                    86 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.44;
                                                                                                                                                         Lentivirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chain type from two human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ų,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                          Timilitie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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RESULT 11
TAT_HYLMA
ID TAT_B
AC P0461
DT 13-AU
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TAT_HV1MA
P04613;
13-AUG-1987
13-AUG-1987
16-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                               Cell 46:63-74(1986).

-!- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING
-!- TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AN
TRANS-ACTIVATING RESPONSIVE TRANSFORM AND/OR ELONGATION FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AIDS.
SEQUENCE
                                                                                                                                    the European Bioinformatics Institute. There are no rest use by non-profit institutions as long as its content modified and this Statement is not removed. Usage by an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alizon M., Wain-Hobson S., Montagnier L., "Genetic variability of the AIDS virus: n of two isolates from African patients.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             deficiency syndrome patients.";

Proc. Natl. Acad. Sci. U.S.A. 83:2209-2213(1986).

-!- EUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATIC RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-slb.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
   HIV; K03456; TATSMAL
                              EMBL; X04415; CAA28015.1;
                                                                              entities requires a license agreement (See http://www.isb-sib.or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                               between
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus type 1 (MAL isolate) (HIV-1). Viruses; Retroid viruses; Retroviridae; Lentivirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TAT protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=86245056; PubMed=2424612;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=11697;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Transcription regulation; Activator; RNA-binding; Nuclear protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRINTS; PR00055; HIVTATDOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR001831; HIV_Tat. Pfam; PF00539; Tat; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; M11840; AAA44999.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -!- SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            47
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                                                                                                                                                                                                                                                                                                           SUBCELLULAR LOCATION: NUCLEAR; NUCLEOLAR.
                                                                                                                                                                                                                                                                                                                                   PROMOTER.
SUBUNIT: BINDS CYCLIN T1 (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                            ACTIVATES TRANSCRIPTION INITIATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PROMOTER.
                                                                                                                                                                                                                   SWISS-PROT entry is copyright. It is produced through a collaboration een the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A04017; TNLJ12.
M11840; TAT$PCV12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         YGRKKRRORRAPQGSQTHQVSLSKQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (Transactivating regulatory protein)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ΑA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05, Created)
05, Last sequence update)
40, Last annotation update)
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••
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         72
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                                                                                                                                                                                                                                                                                                                                                                                            ELONGATION FROM THE LTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 86
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                                                                                                                                           and for
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KVIM-HUMAH
ID KVIM-H
ID KVIM-H
ID KVIM-H
ID KVIM-H
ID KVIM-H
AC P01605
AC P0
RESULT 13
KV10_HUMAN
ID KV10_HUMAN
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Best Local S
Matches 12
                                                                                                                                                                                                        Query
Best 1
                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Scand. J. Immunol. 5:577-684(1976).

-I- MISCELLANEOUS: THE SECOND AND THIRD HYPERVARIABLE REGIONS OF THIS CHAIN ARE IDENTICAL WITH THOSE OF THE HUMAN POM V-III KAPPA CHAIN, WITH WHICH IT SHARES CETTAIN IDIOTYPIC DETERMINANTS.
-I- MISCELLANEOUS: THIS CHAIN WAS ISOLATED FROM AN IGM WITH ANTI-GAMMA GLOBULIN ACTIVITY.
PIR: A01871; KIHULY.
HSSP: P01607: IREI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-JUL-1986
21-JUL-1986
15-JUL-1999
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                        DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Complete amino acid sequence of the variable domains of IgM anti-gamma globulins (Lay/Pom) with shared idiotypic specificities.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                               DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR003006; Ig_MH
InterPro; IPR003596; Ig_v.
Pfam; PF00047; Ig; 1.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Capra J.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KV1M_HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                       DOMAIN
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-JUL-1986 (Rel. 01, Last sequence update)
-JUL-1999 (Rel. 38, Last annotation update)
kappa chain V-I region Lay.
                                                                                                                                                                                                                             Match
                                                                                                                                              SGSGTDATAPASSLQAED 29
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                                                                                                        SGSGTDFTFTISSLQPED
                                                                                                                                                                                     Similarity
13; Conser
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v region.
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        STANDARD
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11834
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72.28;
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0; M1
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                                                                                                                                                                                                                                                                                                                         COMPLEMENTARITY - DETERMINING - 3 FRAMEWORK - 4.
BY SIMILARITY.
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Pred.
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COMPLEMENTARITY-DETERMINING-1
FRAMEWORK-2.
                                                                                                                                                                                                                         Score
                                                                                                                                                                                                                                                                                                                                                                                  FRAMEWORK - 3
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          PRT;
                                                                                                                                                                                                                                                                                    739993A95431434A CRC64;
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No.
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                                                                                                                                                                                                                             Length 108
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Qy

12

SGSGTDATAPASSLQAED

Query Match 39.
Best Local Similarity 72.
Matches 13; Conservative

. 28

Score 57; DB 1 Pred. No. 0.49;

Length 108;

0

Mismatches

Indels

0; Gaps

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MEDIJINE-76039968; PubMed=1182131;

Epp O., Lattman E.E., Schiffer M., Huber R., Palm W.;

The molecular structure of a dimer composed of the variable portions of the Bence-Jones protein REI refined at 2.0-A resolution.";

Biochemistry 14:4943-4952(1975).

-I- MISCELLANEOUS: THE C REGION OF THIS CHAIN HAS THE INV (1,2)
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21-JUL-1986
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and characterization of the tryptic peptides; the complete amino
sequence of the protein; a contribution to the elucidation of the
three-dimensional structure of antibodies, in particular their
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                                                                                                                                                                                                                                                                                                                                                                                 combining site.";
                                                                                                                                                                                                                                                                                                                                                                                                                         Palm W., Hilschmann N.;
"The primary structure of a crystalline monoclonal immunoglobuling
                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-76023758; PubMed-809329;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (Human)
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1REI; 17-FEB-84.
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COMPLEMENTARITY-DETERMINING-2.
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9E8143E1188BCE2A
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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01-NOV-1995 (Rel. 32, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
19 kappa chain V-I region WAT.
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"Characterization and preliminary crystallographic data on the VL-
"Characterization and preliminary crystallographic data on the VL-
related fragment of the human kI Bence Jones protein Wat.";
J. Wol. Biol. 147:185-193(1981).
-i- MISCELAMEOUS: THIS IS A BENCE-JONES PROTEIN.

PDB: 1WTL; 01-NOV-94.
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MEDLINE-81267384; PubMed=6167731;
Stevens F.J., Westholm F.A., Panagiotopoulos N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Solomon A., Stevens F.J., Schiffer M.; "Comparison of crystal structures of two homologous proteins: structural origin of altered domain interactions in immunoglobulin light-chain dimers.";
                                      Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
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SMART; SM00406; IGV; 1.
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FRAMEWORK-2.
COMPLEMENTARITY-DETERMINING-2.
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Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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Braun M., Leibold W., Barnikol H.U., Hilschmann N.;
"Principle of antibody structure. The primary structure of a
monoclonal kappa I-type immunoglobulin L-chain (Bence Jones protein
Bi). 3. The complete aminno acid sequence and the genetic
significance of the variability principles for the mechanism of
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HSSP; P01607; 1REI.
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COMPLEMENTARITY-DETERMINING-3.
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071973
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                              O9fty9 oryza sativ
O71973 human immun
O40231 human immun
O40234 homo sapien
O9n749 homo sapien
O9n749 homo sapien
O71968 human immun
O90217 human immun
O90217 human immun
P90117 human immun
P90118 human immun
P90118 human immun
O00m88 human immun
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45	44	4 4	41		39	38	37	36	3 5	34	w	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17
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ALIGNMENTS

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    Query Match
Best Local Similarity
Matches 13; Conserv
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PRINTS; PR00109; TYKKNASE.

SMART; SM00210; TYKC; 1.

SMART; SM00219; TYTKC; 1.

PROSTIE; PS50011; PROTEIN_KINASE_DOM; 1.

PROSTIE; PS00109; PROTEIN_KINASE_TYR; 1.

ATP-binding; Kinase; Receptor; Transferase.

SEQUENCE 399 AA; 45329 MW; E2D92FEIF6DAAC7B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Oryza sativa nipponbarė(GA3) genomic DNA, chromosome 1, clone:P0456604.";
Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases.
EMBL, AP002818; BAB16226.1;
InterPro; IPR000719; Euk_pkinase.
InterPro; IPR000719; Esr_thr_pkinase.
InterPro; IPR001245; Tyr_pkinase.
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Q9FTY9;
01-MAR-2001 (TIEMBLITEL 16, Created)
01-MAR-2001 (TIEMBLITEL 16, Last sequence update)
01-DEC-2001 (TIEMBLITEL 19, Last annotation update)
PUTATIVE RECEPTOR SER/THR PROTEIN KINASE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota: Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
MCBI_TaxID=4530;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN-CV. NIPPONBARE;
Sasaki T., Matsumoto T., Yamamoto K.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oryza sativa (Rice).
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        Conservative
                                       45.18;
    ;
    Score 65; DB
Pred. No. 0.31
1; Mismatches
                                           DB 10; Length 399; 0.31;
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Q9nea9 leishmania Q9vdv0 drosophila P90148 human imm

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Best Local
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SEQUENCE .....
STRAIN-HE93-A3TAT;
STRAIN-HE93-A3TAT;
MEDLINE-97335179; PubMed-9191845;
Kirchhoff F., Greenough T.C., Hamacher M., Sullivan J.L.,
Escrosiers R.C.;
*Activity of human immunodeficiency virus type 1 promoter/TAR regions and tatl genes derived from individuals with different rates of morraession.";
                                                                                                                                                                                                                                                                                                                                                                              040231;
040231;
01-JAN-1998
01-JAN-1998
01-DEC-2001
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01-AUG-1998
01-AUG-1998
01-DEC-2001
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EMBL; AF049495; AAC68854.1; -.
INTERPO; IPRO01831; HIV_Tat.
Pfam; PF00539; Tat; 1.
PRINTS; PR00055; HIVTATDOMAIN.
ACTLYATO: Nuclear protein; RNA-binding; Transcription regulation
ACTLYATO: Nuclear protein; RNA-DINDINGS CRC64;
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Submilted (FEB-1998) to the EMBL/GenBank/DDBJ databases.

-I-FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE L'
                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                        Viruses; Retroid
NCBI_TaxID=11676;
                                                                                                                                                                                                                                                                                 Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae;
                                                                                                                                                                                                                                                                                                                                                              TAT PROTEIN
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"Molecular cloning and characterization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE-98440562; PubMed-9765443;
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(TrEMBLrel. 19,
(TRANSACTIVATING
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Last annotation updat
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Best Local 9
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Best Local
                                                                                                      Q9NR59;
01-OCT-2000
01-OCT-2000
01-DEC-2001
                                                                              01-DEC-2001 (TIEMBLrel. 15, Last sequence update)
SER/ARG-RICH PRE-MRNA SPLICING FACTOR SR-A1.
SR-A1.
                                                                                                                                                                                                                                                                                                                    Submitted (MAR-1997) to the EMBL/GenBank/DDBJ databases EMBL; AB002405; BAA24179.2; -.
InterPro; IPRR0005115; BPD_TRANSP_INN_MEMBR; UNKNOWN_1.
SEQUENCE 444 AA; 50427 MW; DD4053A374A00FA1 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         O43284;
01-JUN-1998 (TrEMBLrel. 06,
01-MAY-2000 (TrEMBLrel. 13,
01-JUN-2001 (TrEMBLrel. 17,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Virology 232:319-331(1997).

1- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS. ACTIVATING RESPONSIVE SEQUENCE (TAR) NA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
                                           Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE-LYMPHOID;
Abe Y., Takaoka Y.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
                                                                                                                                                       Q9NR59
                                                                                                                                                                                                                                                                                                                                                                                  Abe Y., Takaoka Y.;
"LAK-4 clone from the membrane lymphotoxin expressing subtraction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         043284
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           SEQUENCE
                                  NCBI_TaxID=9606;
                                                                    Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (Human).
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InterPro; IPR001831; HIV_Tat.
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                                                                                                                                                                                                                                                                        Local Similarity
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PROMOTER (BY SIMILARITY).
PROMOTER (BY SIMILARITY).
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Kyriakopoulou
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78.9%;
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5; Mismatches
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Pred. No.
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                                             Craniata; Vertebrata;
Catarrhini; Hominidae
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Catarrhini; Hominidae
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                                               Hominidae;
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 Diamandis
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                                                          Euteleostom1;
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 E.P.;
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071968;
01-AUG-1998
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01-DEC-2001
TAT PROTEIN
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09H7N4;
01-MAR-20Q1 (TEMBLrel. 16,
01-MAR-2001 (TEMBLrel. 16,
01-DEC-2001 (TEMBLrel. 19,
MEDLINE=98440562; PubMed=9765443; Mwaengo D.M., Novembre F.J.; "Molecular cloning and characterization chimpanzees with pathogenic human immunc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRINTS;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Cloning of a gene (SR-A1) encoding a new member of the rich family of pre-mRNA splicing factors."; Submitted (APR-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                               SEQUENCE FROM N.A STRAIN-HIV-1JC;
                                                                                                                                                                                                                  NCBI_TaxID=11676;
                                                                                                                                                                                                                                                                           Human immunodeficiency virus type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
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InterPro; IPR002965; P_rich_extensn
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Submitted (AUG-2000)
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Mammalia; Eutheria;
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07,
                                                                                                                                                                                                                                              Retroviridae;
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AIDS 0:0-0(2001).
AIDS 0:0-0(2001).
                                                                                                                                                                                                                                                                                        Q69622;
Q69622;
01-NOV-1996
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Viruses;
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01-DEC-2001
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EMBL; AF049494; AAC68845.1; -
IntorPro; IPR001831; HIV_Tat.

Pfam; PF00539; Tat; 1.

PRINTS; PR00539; Tat; 1.

PRINTS; PR0055; HIVTATDOMAIN.

ACTIVATOF; Nuclear protein; RNA-binding; Transcription regulation regulation in the companion of the co
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STRAIN+HIV-1JG;

STRAIN-HIV-1JG;

MWaengo D.M., Novembre F.J.;

Submitted (FEB-1998) to the EMBL/GenBank/DDBJ databases.

-I- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE
TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND

TRANS-ACTIVATING RESPONSIVE SEQUENCE SEQUENCE RESPONSIVE SEQUENCE RESPONSIVE
                                                   Human immunodeficiency virus
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J. Virol. 72:8976~8987(1998).
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16; Conserv
     Retroid viruses;
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6 (TrEMBLrel. 01, I
1 (TrEMBLrel. 19, I
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AA; 11572 MW;
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     Retroviridae;
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                                                                                                                                            REGULATORY PROTEIN) (FRAGMENT).
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3; Mismatches
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J. Virol. 69:3273-3281(1995).

-I- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=95264414; PubMed=7745674; Diaz R.S., Sabino E.C., Mayer A., Mosley "Dual human immunodeficiency virus type
                                                                                                                                                                                                                                                                                                                                                                                                                                             ACTIVATES TRANSCRIPTION INITIATION AND/OR ELONGATION PROMOTER (BY SIMILARITY).
EMBL; U80463; AAB39113.1; ...
Interpro; IPR001831; HIV_Tat.
Pfam; PF00539; Tat; 1.
PRINTS; PR00055; HIVTATDOMAIN.
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Quinones 'Mateu M.E., Domingo E.;
Quinones 'Mateu M.E., Domingo E.;
Polnt mutant frequency and intrasubtype B recombination lead
"Polnt mutant frequency and intrasubtype B recombination lead
Venezuelan HIV-1 genetic diversities.";
Venezuelan HIV-1 genetic diversities.";
Submitted (NOV-1996) to the EMEL/GenBank/DDBJ databases.
Submitted (NOV-1996) to the EMEL/GenBank/DDBJ databases.

1-- FUNCTION: TRANSCRIPTIONAL REGULATOR THAT ACTS BY BINDING TO THE
TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND
TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND
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EMBL; Ull185; AAA78871.1; -.
InterPro; IPR001831; HIV_Tat.
                                                                                                                                                                                                                                                                                                                                       Activator; Nuclear protein; RNA-binding; Transcription regulation NON_TER 72 72 72 SEQUENCE 72 AA; 8214 MW; F79D7B8D0205EEB0 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus type 1. Viruses; Retroid viruses; Retroviridae;
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(TrEMBLrel. 03, Last sequence update)
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(TRANSACTIVATING REGULATORY PROTEIN) (FRAGMENT).
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P90128;
01-MAY-1997
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"Point mutant frequency and intrasubtype B recombination lead

"Point mutant frequency and intrasubtype B recombination interaction in Education That acts by Binding To The

TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND

ACTIVATIES TRANSCRIPTION INITIATION AND/OR ELONGATION FROM THE LTR

PROMOTER (BY SIMILARITY).

EMBL; U80477; AAB39127.1; -
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Quinones-Mateu M.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae;
Carr J.K., Avila M., Gomez-Carrillo M., Salomon H Watanaveeradej V., Pando M., Negrete M., Russell Birx D., Andrade R., Vinoles J., McCutchan F.; Submitted (MAY-2001) to the EMBL/GenBank/DDBJ dat EMBL; AF385936; AAK69333.1; -. SEQUENCE 101 AA; 11416 MW; 360605029A87A327 C
                                                                                                                                                                                                                                                                                                                                                                                                                                               Carr J.K., Avila M., Gomez-Carrillo M., Salomon H.,
Watanaveeradej V., Pando M., Negrete M., Russell K.,
Birx D., Andrade R., Vinoles J., McCutchan F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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NON_TER 72 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR001831; HIV_Tat. Pfam; PF00539; Tat; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=11676;
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Viruses; Retroid viruse
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                                                                                                                                                                                                                         STRAIN-ARMA159;
                                                                                                                                                                                                                                                                                                                                    "Diverse BF recombinants have spread widely since the introduction HIV-1 into South America."; AIDS 0:0-0(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN-ARMA159;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TAT PROTEIN
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7 (TrEMBLTel. 03, L
1 (TrEMBLTel. 19, L
V (TRANSACTIVATING)
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                                                                                                                                                                                                                                                                 N.A.
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es; Retroviridae;
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L K., Sanchez J.
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01-MAY-2000 (TREMBLIEL 1
01-MAY-2000 (TREMBLIEL 1
01-JUN-2001 (TREMBLIEL 1
CG6184 PROTEIN.
Drosophila melanogaster (Fruit fly).
Bukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
NCBI_TaxID-7227;
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Smith D.F.
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PROSITE; PS50067; KINESIN_MOTOR_DOMAIN2; 1.
ATP-binding; Colled coll; Microtubules; Motor protein.
NON_TER 829 829
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InterPro; IPR001752; Kinesin.
Pfam; PF00225; Kinesin; 1.
Pfam; PF00380; KINESINHEAVY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=98146435; PubMed=9477341;
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Eukaryota; Euglen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q9NEA9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    294
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 ERKKRRQRRRSGSGTDATAPASSLQAE 28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EQRRLRQRRRGGKRADATAVMMQAQQE 320
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13; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ) (Tremblrel.
) (Tremblrel.
! (Tremblrel.
LATED PROTEIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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48.1%;
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15, Last sequence update)
19, Last annotation update)
(FRAGMENT).
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Last annotation update)
                                                                                                                                                                                                                                                                                                                        Created)
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Pred. No.
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Pred. No. 0.38
3; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31B286D811B7D648 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Quail M.,
                                                                                                                                                                                                                                                                                                                                                                                      989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       829
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zhang L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 5; Length 829;
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RA Ballew R. M. Basu A. Baxendale J. Bayraktaroglu L. Beasley E.M.,
RA Beeson K.Y. Benos P.V. Berman B.P. Bhandari D. Bolshakov S.R.
Berkova D. Bottchan M.R. Bouck J. Brokstein P., Brottier P.,
RA Borkova D. Bottchan M.R. Bouck J. Brokstein P., Brottier P.,
RA Cherry J.M. Cawley S. Dahlke C. Davenport L.B. Davies P.,
RA Cherry J.M. Cawley S. Dahlke C. Davenport L.B. Davies P.,
RA Dodson K. Doup L.E. Downes M. Dugan-Rocha S. Dunkov B.C. Dunn P.,
RA Durbin K.J. Evangelista C.C., Ferricra S., Fleischmann W.,
RA Glodek A. Gong F. Gorrell J.H. Gu Z. Guan P. Harris M.,
RA Harris N.L. Harvey D. Hoiman T.J. Hernandez J.R. Houck J.,
RA Harris N.L. Harvey D. Hoiman T.J. Wei M.-H. Ibegwam C.
RA Liu X. Mattel B. McIntosh T.C. McLeod M.P. Harris M. X.
RA Kimmel B.E. Kodira C.D., Kraft C., Kravitz S., Kulp D. Lai Z.,
RA Liu X. Mattel B. McIntosh T.C. McLeod M.P. McPherson D.,
RA Merkilov G., Milshina N.V., Mobarry C., Morris J., Mesharfi A.,
RA Melson D.R., Nelson K.A. Nixon K. Nusskern D.R. Pacleb J.M.,
RA Rainert K., Remington K., Sanders R.D.C., Scheeler F., Shen H.,
RA Rainert K., Spradling A.C., Stapleton M., Skupski M.P., Smith T.,
RA Rainert K., Remington K., Sanders R.D.C., Scheeler F., Shen H.,
RA Syler E., Spradling A.C., Stapleton M., Skupski M.P., Smith T.,
RA Shue B.C., Siden Kiamos I., Simpson M., Skupski M.P., Smith T.,
RA Mang Z.-Y. Wassarman D.A., Weinstock G.M., Weissenbach J. R.
RA Mang Z.-Y. Wassarman D.A., Weinstock G.M., Weissenbach J. R.
RA Mang Z.-Y. Wassarman D.A., Weinstock G.M., Weissenbach J. R.
RA Mang Z.-Y. Wassarman D.A., Weinstock G.M., Weissenbach J. R.
RA Mang Z.-Y. Wassarman D.A., Weinstock G.M., Weissenbach J. R.
RA Mang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J. R.
RA Globs R. A., Myers E.M., Rubin G.M., Venter J.C., Shu X., Smith H.O.,
RA Gibbs R. A., Myers E.M., Rubin G.M., Venter J.C., Shu X., Smith H.O.,
RA Gibbs R. A., Myers E.M., Rubin G.M., Venter J.C.,
RA Gibbs R. A., Shu S., Shu B., Shu B.,
                                                                                                                                                                                                                                                                                                   P90148
               RRRROCS COLUMN
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Best Local Similarity
                                                                                                                                                                                              01-MAY-1997
01-MAY-1997
01-DEC-2001
SEQUENCE FROM N.A.
STRAIN-PATIENT LTSS;
MEDLINE-99049627; Pubmed-9833881;
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Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F.,
George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N.,
Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfelffer B.D.,
Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.,
Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D.,
                                                                                         NCBI_TaxID=11676;
                                                                                                                                  Human immunodeficiency
                                                                                                                                                                                                                                                             P90148
                                                                                                                                                                                                                                                                               P90148
                                                                                                                                                                            TAT PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A. STRAIN-BERKELEY;
                                                                                                                                                                                                                                                                                                                                                                                 816 KOKKRRKRRESASGITTTASEDSAGVEE
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                                                                                                                  Retroid
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                                                                                                                                                                            (TRANSACTIVATING
                                                                                                                                                                                                                                                                               PRELIMINARY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                  viruses;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    41.78;
                                                                                                           virus type 1
es; Retroviridae;
                                                                                                                                                                 Last sequence update)
Last annotation update)
G REGULATORY PROTEIN) (FRAGMENT).
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                                                                                                                                                                                                                                                                                                                                                                                    843
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                                                                                                                                                                                                                                                                          72
                                                                                                             Lentivirus
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Search completed: July 15, 2002, 13:22:48 Job time: 1484 sec
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RA Lederman M.M., Domingo E.;

RT "LTR and tat variability of HIV-1 isolates from patients with

RT divergent rates of disease progression.";

RI Virus Res. 57:11-20(1998).

CC TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND

CC TRANS-ACTIVATING RESPONSIVE SEQUENCE (TAR) RNA ELEMENT AND

CC PROMOTER (BY SIMILARITY).

DR EMBL; U80544; AAGB2983.1; ".

DR InterPro; IPRO01831; HIV_Tat.

DR Pfam; pF00539; Tat. 1.

DR Pfam; pF00539; Tat. 1.

DR PRINTS: PR00055; HIVTANDOMAIN.

KW ACTIVATOR PRO01831; HIVTANDOMAIN.

KW ACTIVATOR; NGLEAR PROCEAR PROMOTER 72

SO SEQUENCE 72 AA; 8340 MW; FED4E3EEE915A6E3 CRC64;
                                                                                                                                                                                                                                                                                      Query Match 41.0%; Score 59; DB 15; Length 72; Best Local Similarity 50.0%; Pred. No. 0.38; Matches 13; Conservative 3; Mismatches 10; Indels
                                                                                                                                                                   1 YERKKRRQRRRSGSGTDATAPASSLQ 26
| ||||||||| |: |: | |
47 YGRKKRRQRRRAPQGSQTNQVSLSKQ 72
                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                           Gaps
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Database
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence:
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seq length: 2000000000
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11:
12:
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16:
1: //SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1980.DAT: *
2: //SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1981.DAT: *
3: //SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1982.DAT: *
3: //SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1983.DAT: *
4: //SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1983.DAT: *
5: //SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1984.DAT: *
5: //SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1985.DAT: *
5: //SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1986.DAT: *
7: //SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1989.DAT: *
7: //SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1989.DAT: *
9: //SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1989.DAT: *
9: //SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1989.DAT: *
7: //SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA1999.DAT: *
7: //SIDS1/gcgdata/hold-geneseq/geneseqp-embl/AA2000.DAT: *
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14.104 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  YERKKRRQRRRSGSGTDATAPASSLQAED
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

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AAB18861	AAB18857	AAW4 24 66	AAR98489	AAR50314	AAR50313	AAW27527	AAR52059	AAY05269	AAW62808	AAW62807	ID	SOMMANTES
Amino acid sequenc	Amino acid sequenc	Interleukin-5 huma	Humanised 2B6 anti	Humanised light ch	Humanised light ch	Light chain variab	Light chain variab	Antibody 24-31 hum	Amino acid sequenc	Amino acid sequenc	Description	

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ALIGNMENTS

RESULT

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AAW62807
   Green L,
Mendez M;
 New transgenic non-human
                    WPI; 1998-333314/29
                                                                                    03-DEC-1996;
                                                                                                                                                                                    Human; immunoglobulin; Ig; transgenic; non-human mammal; inactivated endogenous Ig locus; B-cell development; human heavy chain Ig locus; micro constant region; J-H; D-H; V-H gene; kappa light chain Ig locus; kappa constant region; J-kappa gene; V-kappa;
                                                                                                      03-DEC-1997;
                                                                                                                         11-JUN-1998
                                                                                                                                           W09824893-A2
                                                                                                                                                                                                                                                          23-SEP-1998
                                                                                                                                                                                                                                                                            AAW62807;
                                                                                                                                                                                                                                                                                            AAW62807 standard; Peptide;
                                                                (ABGE-) ABGENIX INC.
                                                                                                                                                              Homo sapiens
                                                                                                                                                                                production; antibody.
                                                                                                                                                                                                                                   Amino acid sequence of a human antibody fragment.
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                                                Jakobovits
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified genome also has an inserted human heavy chain Ig locus in migraphile genome also has an inserted human heavy chain Ig locus in micro constant region and regulatory and switch sequences, human J-H genes, human D-H genes, and human v-H genes and an inserted human kappa light chain Ig locus in germline configuration, the human kappa light chain Ig locus comprising a human kappa constant region, J-kappa genes, and v-kappa genes, where the number of v-H and v-kappa genes inserted are selected to restore normal B-cell development in the mammal. The transgenic animals have a near complete human Ig locus, including both a human heavy chain locus and a human kappa light chain locus. They can be used for the production of human antibodies when exposed to particular antigens e.g. when exposed to human II-B, EGFR or TNF- alpha the mice will produce antibodies to II-B, EGFR or TNF- alpha
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunoglobulin locus and a near complete human immunoglobulin locus, used for production of human antibodies
  New transgenic non-human mammals - having an inactivated immunoglobulin locus and a near complete human immunoglo used for production of human antibodies
                                                                                                                                                                                                                    03-DEC-1996;
                                                                                                                                                                                                                                                       03-DEC-1997;
                                                                                                                                                                                                                                                                                                                                   W09824893-A2
                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                           inactivated endogenous Ig locus; B-cell development;
human heavy chain Ig locus; micro constant region; J-H; D-H; V-H gene;
kappa light chain Ig locus; kappa constant region; J-kappa gene; V-kappa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; immunoglobulin; Ig; transgenic; non-human mammal; inactivated endogenous Ig locus; B-cell development;
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                                                                                                                                                                            (ABGE-) ABGENIX INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              acid sequence of a human antibody fragment.
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Pred. No.
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                       immunoglobulin locus
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         specification describes a transgenic non-human mammal which has genome modifications that comprise an inactivated endogenous immunoglobulin (Ig) locus, so that the mammal does not display normal B-cell development. The modified genome also has an inserted human heavy chain Ig locus in germline configuration, the human heavy chain Ig locus comprising a human micro constant region and regulatory and switch sequences, human J-H
                                                                                                                                                                                                                                                                                                                                                                    Antibody 24-31; variable light chain; humanised antibody; gene therapy; CD4() binding inhibitor; gp39; autoimmune disease; food-related allergy; graft-versus-host disease; graft rejection; intestinal inflammation; reversible obstructive airways disease; allergy; transplant rejection; asthma; leukaemia; lymphoma; immunosuppressant; cellular therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 79; 128pp; English
 Claim 7; Page 84; 122pp; English
                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                       Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antibody
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                                  New humanized antibodies
                                                                                             Black A,
                                                                                                                                                           08-SEP-1997;
                                                                                                                                                                                            08-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                      immune response
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                                                                                                                                                                                                                                                                                                      sapiens
                                                                                                                              IDEC PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-31 humanised VL chain sequence.
                                                                                               Hanna N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94
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                                                                                                                                                                                            98WO-US18163
                                                                                                                                                                                                                                                                                                                                                      inhibitor; cancer
                                                                                                Newman
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                                  to human
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Pred. No. 0.24;
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RESULTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antibody; humanised; murine; human; heavy chain; light; variable; framework region; complementarity determining region; reshaping; modelling; surface residue; modify.
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   09-SEP-1992;
                                                               07-SEP-1993;
                                                                                                                        13-APR-1994
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                                                               93EP-0307051
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24..40
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96..104
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63..95
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /label= complementarity_determining_region_l
/note= "CDR 1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /label- framework_region_1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      43.88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "CDR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "FR 1"
                                                                                                                                                                                                                                                                                                             "FR 3"
                                                                                                                                                                                                                                                                                                                                                                         "CDR 2"
                                                                                                                                                                                                                                                                                                                                                                                                                                     "FR 2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of human KV4B antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 63;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 107;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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AAW27527
ID AAW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Modification of a rodent antibody (Ab) or fragment by resurfacing in order to produce a humanised rodent Ab can be determined by calculating homology between murine and human Ab antibody surfaces. In order to test the resurfacing approach of the invention, three humanisation experiments were set up: (1) traditional loop graffing; (2) resurfacing approach using most similar chain; and (3) resurfacing approach using human sequences with most similar surface residues. The Ab used was the murine anti-N901 Ab (see AAB2005). Experiment 3 was carried out using the present sequence which represents the human KV4B ab light chain variable region with 71 percent homology with anti-N901 Ab. N901/KV4B (AAR52060) was prepd. by resurfacing.
                                                                                       (ASAK ) ASAHI
(NIKK-) NIKKA
(TORI ) TORII
(TSUR/) TSURA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Method of resurfacing of rodent antibodies to produce humanised antibody forms - for producing non-human antibodies with improved therapeutic efficiency by presenting human surface on V-region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Fig 4A; 230pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Guild BC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PEDE/) PEDERSEN J T. (IMMU-) IMMUNOGEN INC
Humanised, semi-chimeric and chimeric antibodies against human high-affinity IgE receptor – useful medicinally and have low antigenicity in humans
                                                                                                                                                                                                                                                                                                                              Complementarity determining region; CDR; murine; mouse; human; high affinity; immunoglobulin E; receptor; monoclonal antibody; IgE; MAD; llght chain; variable region; humanised; semi-chimeric;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1994-120230/15
                                                                                                                                                                                                                                                       JP09191886-A
                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                 chimeric; treatment;
                                                                                                                                                                                                                                                                                                                                                                                       Light chain variable region
                                                                                                                                                                                                                                                                                                                                                                                                                     16-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW27527;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW27527 standard; Protein; 112 AA
                                                            WPI; 1997-429186/40.
                                                                                                                                                                 19-JAN-1996;
                                                                                                                                                                                              19-JAN-1996;
                                                                                                                                                                                                                          29-JUL-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        111 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                      WHISKEY
                                                                                                                                    BREWERIES LTD
                                                                                                       YAKUHIN
                                                                                                                                                                 96JP-0024816
                                                                                                                                                                                              96JP-0024816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             43.88;
77.88;
                                                                                                                                                                                                                                                                                                               prevention; disease; allergy; CRA2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rees AR,
                                                                                                                                                                                                                                                                                                                                                                                       of human CRA2 antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 63; I
Pred. No. 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Roguska MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 15; Length 111; 0.28;
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Best Local
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                                                                                                     Charoenvit Sadoff JC,
                                    New engineered antibodies and fusion proteins plasmodium infection - contg. murine antibody corresp. nucleic acid, vectors and transformed
Naturally-occurring
                   Claim 5;
                                                                                                                                          (SMIK )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence, the light chain variable region of the human antibody (Ab) CRA2, was used in the preparation of a humanised or semi-chimeric monoclonal Ab (MAb), comprising complementarity determining regions (CDR) from a murine, anti-human high affinity immunoglobulin E (9g) receptor, MAb. The humanised, semi-chimeric or chimeric MAb can be used to treat or prevent diseases, specifically allergies, associated with the receptor, and has very
                                                                                      WPI;
                                                                                                                                                                       09-SEP-1992;
                                                                                                                                                                                          08-SEP-1993;
                                                                                                                                                                                                             17-MAR-1994
                                                                                                                                                                                                                               WO9405690-A
                                                                                                                                                                                                                                                           Region
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                                                                                                                                                                                                                                                                                                Region
                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                            complementarity determining region; fusion protein; murine; variable; light; heavy; chain; malaria.
                                                                                                                                                                                                                                                                                                                                                               Monoclonal antibody; Plasmodium falciparum;
                                                                                                                                                                                                                                                                                                                                                                                Humanised light chain variable region Pfhzlc1-1
                                                                                                                                                                                                                                                                                                                                                                                                      05-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                        AAR50313
                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR50313 standard; Protein; 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                   (USNA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12 SGSGTDATAPASSLQAED 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                             o,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antigenicity in humans.
                                                                                    1994-101115/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sgsgtdftltisslgaed 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14;
                                                                            AAQ44827
                                                                                                                                 SMITHKLINE BEECHAM
US SEC OF ARMY.
US SEC OF NAVY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                   Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 14; 26pp;
                                                                                                     Y, Hoffman
Sylvester I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            112 AA;
                   ຸນ
                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                       92US-0941654
                                                                                                                                                                                         93WO-US08435
                 98pp;
                                                                                                                                                                                                                                                           /label= CDR2
95..103
                                                                                                                                                                                                                                                                             /label= CDR1
56..62
                                                                                                                                                                                                                                                  /label=
                                                                                                                                                                                                                                                                                                       Location/Qualifiers
(AAQ44841-42) and
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                   English
                                                                                                                                                                                                                                                  CDR3
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                                                                                                     Hurle
Gross
                                                                                                                                                    CORP
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Pred.
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                                                                                                     X X
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                                     transformed
                                                                                                              Rosenberg
synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 18;
0.28;
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                                    s for preventing
y CDR sequences,
ed cells
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(AAQ44825-28)
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                                              and
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RESULT
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Best Local
     Naturally-occurring (AAQ44841-42) and synthetic (AAQ44825-28) variable light chain and variable heavy chain sequences derived from murine mAb NFS2 are provided. Murine mAb NFS2, its variable chain peptides, CDMs, functional fragments, Fab fragments, and analogs are useful in prodn. of fusion
                                                                                                                                               Claim
                                                                                  New engineered antibodies and fusion proteins 
Plasmodium infection - contg. murine antibody of 
corresp. nucleic acid, vectors and transformed
                                                                                                                                                                                               (SMIK )
                                                                                                                           N-PSDB;
                                                                                                                                     WPI; 1994-101115/12.
                                                                                                                                                                                     (USNA
                                                                                                                                                                                                                           09-SEP-1992;
                                                                                                                                                                                                                                                08-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                      Monoclonal antibody; Plasmodium falciparum; CDR; complementarity determining region; fusion prote murine; variable; light; heavy; chain; malaria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            variable light chain and variable heavy chain sequences derived from murine mAb NFS2 are provided. Murine mAb NFS2, its variable chain peptides, CDRs, functional fragments, Fab fragments, and analogs are useful in prodn. of fusion proteins, esp. engineered antibodies. These prods. are
                                                                                                                                                                                                                                                                   17-MAR-1994
                                                                                                                                                                                                                                                                                      W09405690-A.
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                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Humanised light chain variable region
                                                                                                                                                                                                                                                                                                                                                          Region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   used to protect humans against Plasmodium infections.
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                                                                                                                                                                                    SMITHKLINE BEECHAM
US SEC OF ARMY.
US SEC OF NAVY.
                                                                                                                           AAQ44828
                                                               Fig
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                  Hoffman
                                                              98pp;
                                                                                                                                                                                                                            9205-0941654
                                                                                                                                                                                                                                               93WO-US08435
                                                                                                                                                                                                                                                                                                                     95
                                                                                                                                                                                                                                                                                                                                      /label= CDR1
56..62
                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers 24.40
                                                                                                                                                                                                                                                                                                                            /label- CDR2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           43.8%;
77.8%;
                                                                English.
                                                                                                                                                       n s,
                                                                                                                                                                                                                                                                                                          CDR3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           88
                                                                                                                                                      Hurle
Gross
                                                                                                                                                                                                        CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           113 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 63;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                        S. Z
                                                                                                                                                                                                                                                                                                                                                                                                                  fusion protein;
                                                                                                                                                                Rosenberg
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                                                                                  CDR sequences,
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engineered

antibodies.

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Best Local S
Matches 14
                                                                                                              06-JUN-1995;
23-DEC-1994;
06-JUN-1995;
                                                                                      (SMIK )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               used
       New monoclonal antibody to human interleukin-5 - used to products for the treatment and diagnosis of conditions as with excess eosinophil prodn., e.g asthma etc.
                                         N-PSDB;
                                                                                                                                               22-DEC-1995;
                                                                                                                                                               11-JUL-1996
                                                                                                                                                                               WO9621000-A2
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                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                      monoclonal antibody; MAb
                                                                                                                                                                                                                                                                                                                                                                           Antibody engineering; humanised antibody; chimeric antibody; Fab; interleukin-5; II-5; eosinophil; asthma; allergic rhinitis; atopic dermattitis; therapy; diagnosis; light chain; VL;
                                                                                                                                                                                                                                                                                                                                                                                                              Humanised 2B6 antibody light chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                               12-OCT-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR98489 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                Holmes
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                                                                        RS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sgsgtdftltisslgaed 88
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                                               1996-333976/33
                                                               SD,
                                                                                      SMITHKLINE BEECHAM CORP
SMITHKLINE BEECHAM PLC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity 77.4
14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protect humans
                                                               Appelbaum ER,
, McMillan LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               113
                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                              95US-0470110.
94US-0363131.
95US-0467420.
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                                                                                                                                               95WO-US17082
                                                                                                                                                                                                                                               /note=
63..94
                                                                                                                                                                                                                                                                                       /note=
41..55
                                                                                                                                                                                                       /note= "mouse
104..113
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                                                                                                                                                                                               /note=
                                                                                                                                                                                                                        /label= CDR3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Protein; 113
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                                                                                                                                                                                                                                       "human
                                                                                                                                                                                                                                                                               "human
                                                                                                                                                                                                                                                                                                                       "human LEN framework region"
                                                                                                                                                                                               "human LEN
                                                                                                                                                                                                                                                                                               "mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              . 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               against Plasmodium infections
                                                                                                                                                                                                                                                       "mouse
                                                               Chaiken IM,
Theisen TW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 63; DB
Pred. No. 0.29
0; Mismatches
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Matches
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                                                                                                                                                                                 Region
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Region
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                                                                                                                                                                                                                                                                                                                                                                                     atopic
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                                                                                                                                                                                                                                                                                                                                                      Chimeric
                                                                                                                                                                                                                                                                                                                                                                                   Neutralising antibody; monoclonal antibody; MAD; 2B6; LEN; mouse; interleukin-5; IL-5; human; eosinophil; asthma; allergic rhinitis; atopic dermatitis; therapy; diagnosis; humanised antibody.
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         21-JUN-1996;
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Pred. No. 0.29;
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Best Local :
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polynúcleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid % \left( 1\right) =\left\{ 1\right\} 
                                                                                                          N-PSDB; AAA96134
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rheumatoid arthritis; coronary heart disease.
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                                                                                                                                                                                                                                                                             (SVIN-) ST VINCENT'S HOSPITAL SYDNEY LTD
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Pred. No. 0.29
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Best Local Similarity
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                                                                                          Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid arthritis and coronary heart disease \,
                                                                                                                                                                                 WPI;
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                                                                                                                                                               N-PSDB; AAA96138.
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The present sequence represents the light chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies

Claim 30; Page 136; 163pp; English

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                                                                                                                                                                                                                                                                                                                                                                          Ward
                                     The present sequence represents the light chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain
  chelators, drugs, prodrugs, or gadolinium. The polypept
                                                                                                                                                                Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antibody;
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                                                                                                                                                                                                          and coronary heart disease
                                                                                                                                                                Page 138;
                                                                                                                                                                                                                                                                                                                                                                          Coomber DWJ;
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polypeptide
                                                                                                                                                           163pp;
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77.88;
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Pred. No.
  components
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RESULT 3
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Best Local
         The present sequence represents the light chain of an antibody reactive against p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies are useful in pharmaceutical compositions, which additionally contain chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in vaccines, for inducing an immune response against a disease in a vertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as well as for gene therapy and recombinant production of the polypeptides. In particular, the following can be treated cancer, rheumatoid arthritis and coronary heart disease. Cancers include carringers tumours, tumours of enthelial origin e.g. cancers include carringers tumours, tumours
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                                                                                                                                                                                                                                                                                           Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid arthritis and coronary heart disease
 of epithelial origin,
                                                                                                                                                                                                                                                      Claim 30; Page 144; 163pp; English.
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77.8%;
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Pred. No. 0.29;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antibody light chain clone 163.15.
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                                                                                        The present sequence represents the light chain of an antibody reactive C against p53. The antibody is obtained from a vertebrate host expressing CC an immune response against a naturally occurring disease. The antibodies CC are useful in pharmaceutical compositions, which additionally contain CC chelators, drugs, prodrugs, toxins and imaging markers e.g. radioisotopes CC or gadolinium. The polypeptide components of the antibodies are useful in CC vaccines, for inducing an immune response against a disease in a CC vertebrate, for treatment and/or prophylaxis of disease and for detection CC purposes. The nucleic acid sequences can be used to detect a disease as CC in particular, the following can be treated cancer, rheumatoid arthritis CC and coronary heart disease. Cancers include carcinogenic tumours, tumours CC epithelial origin, e.g. colo-rectal cancer, prostate cancer lung cancer, CC chead and neck tumours, hepatic cancer, pancreatic cancer, ovarian cancer, CC urinary/genital tract cancer, badder cancer, prostate cancer and curinary/genital tract cancer, cossophageal cancer, mesenchymal tumours, and hemopolistic tumours, e.g. a Reference and cancer and cancer and cancer, brain cancer, tumours, e.g. a Reference and cancer and cancer and cancer, brain cancer, cossophageal cancer, mesenchymal tumours,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 30; Page 146; 163pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polynucleotides encoding anti-p53 antibodies, polypeptides and peptide fragments, useful in treatment and diagnosis of cancer, rheumatoid
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Pred. No.
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                                                                                The present sequence represents the light chain of an antibody reactive capainst p53. The antibody is obtained from a vertebrate host expressing an immune response against a naturally occurring disease. The antibodies care useful in pharmaceutical compositions, which additionally contain chelators, drugs, prodrugs, toxins and immaging markers e.g. radioisotopes or gadolinium. The polypeptide components of the antibodies are useful in veccines, for inducing an immune response against a disease in a covertebrate, for treatment and/or prophylaxis of disease and for detection purposes. The nucleic acid sequences can be used to detect a disease as computed as for gene therapy and recombinant production of the polypeptides. In particular, the following can be treated cancer, renumatoid arthritis and coronary heart disease. Cancers include carcinogenic tumours, tumours of epithelial origin, e.g. colo-rectal cancer, prostate cancer of gastric cancer, brain cancer, bladder cancer, prostate cancer and urinary/genital tract cancer, ossophageal cancer, mesenchymal tumours, and hemonietic tumours, e.g. a figel lymbons.
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Sequence
                                                           e.g. sarcoma, and hemopoietic tumours, e.g. B cell lymphoma.
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Copyright (c) 1993 - 2000 Co
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STRANDEDNESS: single; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-554-840-15
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                                                                                                                       GENERAL INFORMATION:
APPLICANT: BLACK, Amelia
APPLICANT: HANNA, Nabil
APPLICANT: PADLAN, Eduaro
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FILING DATE: 07-NOV-1995
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35,030
REFERENCE/DOCKET NUMBER: 012712-127
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-6620
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 80 amino acids
                                 CORRESPONDENCE ADDRESS:
                                            APPLICANT: NEWMAN, ROLAND A.
TITLE OF INVENTION: HUMANIZED ANTIBODIES TO HUMAN 9p39,
TITLE OF INVENTION: COMPOSITIONS CONTAINING AND THERAPH
NUMBER OF SEQUENCES: 28
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
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CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION: COMPOSITIONS CONTAINING AND THERAPEUTIC USE THEREOF
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CITY: Alexandria
STATE: Virginia
                                                                                                                                                                                                                                                                                                            47 SGSGTDFTLTISSLQAED 64
                                                                                                                                                                                                                                                                                                                                           12 SGSGTDATAPASSLQAED 29
                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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               ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Burns, Doane, Swecker & Mathis P.O. Box 1404
P.O.
                                                                                                                       HANNA, Nabil
PADLAN, Eduardo A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            United States
                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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Burns, Doane, Swecker & Mathis
O. Box 1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Roland A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eduardo A.
                                                                                                                                                                                                                                                                                                                                                                                                43.8%;
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                                                                                                                                                                                                                                                                                                                                                                                              Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                  0.03;
                                                                                                                                                                                                                                                                                                                                                                                                                    DB 3;
                                                                    AND THERAPEUTIC USE THEREOF
                                                                                                                                                                                                                                                                                                                                                                                 4
                                                                                                                                                                                                                                                                                                                                                                                                                    Length 80;
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RESULT 4
US-08-525-539A-81
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 81, Application US/08525539A Patent No. 6309636
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: DO COUTO, FERNANDO J.R.
APPLICANT: CERIANI, ROBERTO L.
APPLICANT: PETERSON, JERRY A.
TITLE OF INVENTION: RECOMBINANT PEPTIDES DERIVED FROM THE
TITLE OF INVENTION: MCS ANTI-BA46 ANTIBODY, METHODS OF USE THEREOF, AND
TITLE OF INVENTION: MCTHODS OF HUMANIZING ANTIBODY PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (703) 836-2021 INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                               ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Teskin, Robin L.
REGISTRATION NUMBER: 35,030
REFERENCE/DOCKET NUMBER: 01:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 836-6620
                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 81
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                         ATTORNEY/AGENT INFORMATION:
NAME: DYLAN, TYLER
REGISTRATION NUMBER: 37,
                                                                                                                                                                                  CURRENT APPLICATION DATA:
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ATTORNEY/AGENT INFORMATION:
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CURRENT APPLICATION DATA:
REFERENCE/DOCKET NUMBER: 27
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
                                                                                                                                                                                                                                                                                                                                                 STREET: / J
                                                                                                                  APPLICATION NUMBER: US/08/525,539A FILING DATE: 14-SEP-1995 CLASSIFICATION: 424
                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12 SGSGTDATAPASSLQAED 29
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Local Similarity 77.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: Sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/0 FILING DATE: 07-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Alexandria
STATE: Virginia
                                                                                                                                                                                                                                                                                                                               COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: United ZIP: 22313-1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14;
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                                                           37,612
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RESULT 6
US-08-467-420A-21
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; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-07-942-245-30
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US-07-942-245-30
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                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/07/942,245
FILING DATE: 09-5EP-1992
CLASSIFICATION: 530
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 293-7060
TELEFAX: (202) 293-7860
TELEFAX: (202) 293-7860
TELEFAX: (202) 293-7860
INFORMATION FOR SED ID NO: 30:
SEQUENCE CHARACTERISTICS:
                                                                                                                   Query Match
Best Local Similarity
Matches 14; Conserv
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COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY disk
COMPUTER: HP 9000/700 Workstation
OPERATING SYSTEM: UNIX
SOFTWARE: In house
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 43.8%;
Best Local Similarity 77.8%;
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (415) 494-0792
TELEX: 706141
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: GUILD,
TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 107 amino acids
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CITY: Washington
STATE: D.C.
                                                                                    12 SGSGTDATAPASSLQAED 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   65 SGSGTDFTLTISSLOAED 82
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                                                         SGSGTDFTLTISSLQAED 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/07942245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: Sughrue, Mion, Zinn, Macpeak & Seas 2100 Pensylvania Avenue, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REES,
ROGUSKA,
                                                                                                                     Conservative
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522
                                                                                                                                  43.8%;
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Anthony R.
Anthony R.
Michael A.
Braydon C.
SURFACE RESIDUE VENEERING OF RODENT
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Pred. No.
                                                                                                                                  Score 63; DB 1;
Pred. No. 0.043;
                                                                                                                   Mismatches
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0.041;
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                                                                                                                Gaps
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US-08-470-110A-21

: Sequence 21, Application US/08470110A

: Patent No. 5693323
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Best Local (
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                                                                                                                                                                                                                                  GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                            APPLICANT:
APPLICANT:
APPLICANT:
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TOPOLOGY: line
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: 610 270-5024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US C
FILING DATE: 23-DEC-1994
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APPLICANT: Appelbaum, Edward
APPLICANT: Chaiken, Irwin M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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CORRESPONDENCE ADDRESS:
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APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Recombinant IL5 Antagonists Useful in TITLE OF INVENTION: Treatment of IL5 Mediated Disorders
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         71 SGSGTDFTLTISSLQAED
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RECOMMINICATION TO THE PERSON PROPERTY PROPERT
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CITY: King of Prussia
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE:
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ZIP: 19406-0939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Sutton, Jeffrey A REGISTRATION NUMBER: 34,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  H: 113 amino acids amino acid
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Holmes, Stephen D.
Cook, Richard M.
Gross, Mitchell S.
Holmes, Stephen D.
                                                                                                             Appelbaum, Edward
Chaiken, Irwin M.
                                                                                                                                                                                          Ames, Robert S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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77.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 63; DB 1;
Pred. No. 0.043;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 113;
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APPLICANT:

APPLICANT:

McMillan, Lynette J Theisen, Timothy W.

APPLICANT:

NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:

TITLE OF INVENTION:

Recombinant IL5 Antagonists Useful in Treatment of IL5 Mediated Disorders

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; MOLECULE TYPE: protein US-08-470-110A-21
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                                                                                                                                                                                                                                                                                                                                         Sequence 21, Application US/08667769A Patent No. 5783184
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 610 270-5090
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acid
                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                      APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                      APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50282
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610 270-5024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
CLASSIFICATION: 426
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                               APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinant IL5 Antagonists Useful in
TITLE OF INVENTION: Treatment of IL5 Mediated Disorders
NUMBER OF SEQUENCES: 76
CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKiine Beecham Corp./Corporate
STREET: P.O. Box 1539-UW2220
                                                                                                                                                                                                                                                                                        APPLICANT:
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: SmithKline Beecham Corp./Corporate ADDRESSEE: Intellectual Property STREET: P. O. Box 1539.UW2220 CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                               COUNTRY:
                                                 CITY: King of Prussia
STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                               71
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14;
                19406-0939
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                                              Pennsylvania
                                                                                                                                                                                                                     Cook, Richard M.
Gross, Mitchell S.
Holmes, Stephen D.
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                                                                                                                                                                                                                                                                                        Appelbaum, Edward R.
                                   USA
                                                                                                                                                                                                   McMillan,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              43.8%;
77.8%;
                                                                                                                                                                                                       Lynette J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US/08/470,110A
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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0.043;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 113;
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GENERAL INFORMATION:
APPLICANT: Ames, Robert S.
APPLICANT: Appelbaum, Edward R.
APPLICANT: Chaiken, Irwin M.
APPLICANT: Cook, Richard M.
APPLICANT: Gross, Mitchell S.
APPLICANT: Holmes, Stephen D.
APPLICANT: Holmes, Stephen D.
APPLICANT: Theisen, Timothy W.
APPLICANT: Theisen, Timothy W.
TITLE OF INVENTION: Recombinant IL5 Antagonists Useful in TITLE OF INVENTION: Treatment of IL5 Mediated Disorders NUMBER OF SEQUENCES: 74
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/363131
FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: SUtton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 610-270-5090 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 01 FILING DATE: 06-JUN-1995 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corp./Corporate
ADDRESSEE: Intellectual Property
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: 610-270-5024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 22-DEC-1995 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/17082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: a
                                                                                STATE: F
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/667,769A FILING DATE:
                                                                                                                       STREET: P. O. Box 1539-UW2220
CITY: King of Prussia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          71 SGSGTDFTLTISSLQAED 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 43.8%;
Local Similarity 77.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE:
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                                                                 19406-0939
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                                                                                                   Pennsylvania
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linear
                                                                                    USA
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Pred. No. 0.043;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-637-647-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: SULTON, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P50282
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1070-5024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 610 270-5090 INFORMATION FOR SEQ ID NO: 21:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCE: 50

CORRESPONDENCE ADDRESS:
ADDRESSE: SmithKline Beecham Corp./Corporate
ADDRESSEE: Intellectual Property
STREET: P. O. BOX 1539-UW2220
CITY: King of Prussia
STATE: Pennsylvania
COUNTRY HEAD
                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: McMillan, Lynette J.

APPLICANT: Theisen, Timothy W.

TITLE OF INVENTION: Recombinant IL5 Antagonists Useful in TITLE OF INVENTION: Treatment of IL5 Mediated Disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                      CLASSIFICATION: 536 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
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                                                                                                                               CURRENT APPLICATION DATA:
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SOFTWARE: Patentin Release #1.0
                                                                                     APPLICATION NUMBER: US/08/637,647
FILING DATE: 19-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      71
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                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                 COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/940,371
CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14;
                                                                                                                                                                                                                                                  19406-0939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amino acid
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Gross, Mitchell S.
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77.8%;
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                                     US/08/363,131
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 63; DB 2;
Pred. No. 0.043;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 113;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
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                                                             US-08-525-539A-80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-525-539A-80
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Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 80,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 610 270-5090 INFORMATION FOR SEQ ID NO: 21:
                                                                                                                                       TELEFAX: (415) 494-079:
TELEX: 706141
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/525,539A
FILING DATE: 14-SEP-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: DYLAN, TYLER
REGISTRATION NUMBER: 37,612
                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 81
CORRESPONDENCE ADDRESS:
ADDRESSEE: WORRISON & FOERS
STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS
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TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
                                                                                                                                                                            TELEPHONE: (415) 494-0792
                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 71 SGSGTDFTLTISSLQAED 88
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Sutton, Jeffrey A. REGISTRATION NUMBER: 34,
                                                                                         TYPE: amino acid
STRANDEDNESS: si
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 94304-1018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH:
                                                                               TOPOLOGY:
                                                                                                                            LENGTH:
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6309636
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77.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   METHODS OF HUMANIZING ANTIBODY PEPTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RECOMBINANT PEPTIDES DERIVED FROM THE MC3 ANTI-BA46 ANTIBODY, METHODS OF USE THEREOF, AND
43.8%;
77.8%;
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Score 63;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 63; DB 3; Pred. No. 0.043;
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                                                                                                                                                                                                                                                                                                                                                                        Version #1.30
DB 4;
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              Length 113;
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Query Match Best Local Similarity

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RESULT 13
PCT-US93-08435-8
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                                                                                             Sequence 8, Applicati
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                   Query Match
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   APPLICANT:
APPLICANT:
TITLE OF IN
                                                APPLICANT:
                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
FILING DATE: 09-SEP-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 61
CORRESPONDENCE ADDRESS:
ADDRESSEE: Howson and Howson
STREET: Box 457, 321 Norristown Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: U.S. Government, Secretary of
APPLICANT: the Army
TITLE OF INVENTION: Novel Antibodies for Conferring Passive
TITLE OF INVENTION: Immunity Against Infection by a Pathogen in Man
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Bak, Mary E.
REGISTRATION NUMBER: 31,215
REFERENCE/DOCKET NUMBER: SBI
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version *1.25 CURRENT APPLICATION DATA:
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CITY: Spring House
STATE: PA
COUNTRY: USA
                                                                                                                                                                                           71
                                                                                                                                                                                                                     12 SGSGTDATAPASSLQAED 29
                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
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                                                                                                                                                                                         SGSGTDFTLTISSLOAED 88
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19477
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   T: U. S. Government, Secretary of T: the Army INVENTION: Novel Antibodies for
                                                                                                            Application PC/TUS9308435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application PC/TUS9308435
                                                                                                                                                                                                                                                                                                                                                                                                                                        (215) 540-5818
                                                U. S. Government, Secretary of the Navy
                                                                              SmithKline Beecham, Corporation
                                                                                                                                                                                                                                                     Conservative
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Novel Antibodies for Conferring Passive
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                                                                                                                                                                                                                                                                  Score 63; DB 5;
Pred. No. 0.043;
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                                                                                                            STATE:
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PCT-US95-17082A-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (215) 540-58 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07,
FILLING DATE: 09-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Bak, Mary E.
                                                                                                                                                                                                                                                                                                                                             APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                           APPLICANT:
APPLICANT:
                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                            APPLICANT: APPLICANT:
                                                                                                                                                                                        NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                        APPLICANT: Theisen, Timothy W. TITLE OF INVENTION: Recombinan TITLE OF INVENTION: Treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/08435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: immunity Against Infection by a Pathogen in Man
NUMBER OF SEQUENCES: 61
CORRESPONDENCE ADDRESS:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                  COUNTRY:
                                                                                                                                       ADDRESSEE: Smithkline Beecham Corp./Corporate STREET: P.O. Box 1539-UW2220
CITY: King of Prussia
                                                                                       ZIP: 19406-0939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   71 SGSGTDFTLTISSLQAED 88
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                                                                                                                     Pennsylvania
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Box 457, 321 Norrist
                                                                                                        USA
                                                                                                                                                                                                                                                                                         Cook, Richard M.
Gross, Mitchell S.
Holmes, Stephen D.
                                                                                                                                                                                                                                                                                                                                             Appelbaum, Edward R. Chaiken, Irwin M.
                                                                                                                                                                                                                                                                                                                                                                                Ames, Robert S.
                                                                                                                                                                                                                                                                         McMillan,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   43.88;
                                                                                                                                                                                                                           Treatment of
                                                                                                                                                                                                                                        Recombinant IL5 Antagonists Useful
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                                                                                                                                                                                                                                                                           Lynette J.
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                                                                                                                                                                                                                           IL5 Mediated Disorders
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 113;
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PRIOR APPLICATION DATA:

US 08/470110

CLASSIFICATION:

FILING DATE:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: PCT/US95/17082A

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US-08-360-125-6
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Best Local :
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ETLING DATE: 06-UUN-1995
PRIOR APPLICATION DATA: US 08/363131
APPLICATION NUMBER: US 08/363131
ETLING DATE: 23-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 610-270-5090 INFORMATION FOR SEQ ID NO: 21:
                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 500 kb
                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Wenderoth, Lind & Ponack
STREET: 805 Fifteenth Street, N.W., #700
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Saiko HOSOKAWA
APPLICANT: TOShiaki TAGAWA
APPLICANT: YOKO HIRAKAWA
APPLICANT: NO. 5767246ihiko
APPLICANT: KAZUhiro NAGAIKE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 113 amino acids
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            CLASSIFICATION: 424
PRIOR APPLICATION NUMBER: 07/905,534
PRILING DATE: June 29, 1992
                                                                                                                OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: 610-270-5024
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                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
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TOPOLOGY: 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Sutton, Jeffrey A. REGISTRATION NUMBER: 34,028 REFERENCE/DOCKET NUMBER: P5
                                                                               APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                 CITY: Washington STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            71 SGSGTDFTLTISSLQAED 88
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nes 14; Conserv
                                                                                                                                                                 COMPUTER:
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APPLICATION NUMBER:
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Specifically Binding to Surface Antigen
Cell Membrane
42
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77.8%;
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0; Mismatches
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; RELEVANT RESIDUES IN SEQ
US-08-360-125-6
Search completed: July 15, 2002, 12:59:20 Job time: 391 sec
                                                                                                                    Matches
                                                                                                                               Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Warren M. Cheek, Jr. REGISTRATION NUMBER: 33,367
REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION: TELEPHONE: 202-371-8850
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FRAGMENT TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 114 amino acid
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TISSUE TYPE:
CELL TYPE: H
CELL TYPE: a
CELL LINE:
ORGANELLE:
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LOCATION:
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                                                                                                                                                                                                                                        DOCUMENT NUMBER:
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                                                                                                                                                                                                                                                                                           VOLUME:
                                                                                                                                                                                                                                                                                                        JOURNAL:
                                                                                                                                                                                                                                                                                                                      TITLE:
                                                                                                                                                                                                                                                                                                                                  AUTHORS:
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